

Identification of tissue/cell specific marker genes and
use thereof

5 Cross References to Related Applications

This application claims the priority of US provisional patent application 60/388994, filed June 14, 2002, the disclosure of which is incorporated herein by reference in its entirety.

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Field of the Invention

The present invention relates to a method for the identification of tissue cell specific marker genes, a method for the determination of a disease state or developmental status of cells/tissue as well as to gene expression profiling of cartilage tissue. More specifically, the invention relates to microarrays containing a plurality of selected human chondrocyte specific sequences and their use for classification of cartilage donor tissue or generation of characteristic gene expression profiles of *in vitro* chondrocyte cultures. Such DNA arrays find use as a standard tool of molecular biology research and clinical diagnostics for all cartilaginous or related tissues.

Background of the Invention

Limitation on current microarray technologies

25 DNA array technology, also known as biochip or microarray technology, is currently revolutionizing modern biology. In this technology, a biological sample is applied to a glass slide or chip covered with an array of immobilized DNA probes. Sample nucleic acid complementary to specific probes on the array hybridizes and can be detected with high sensitivity with automated, computerized detectors. In this manner, hundreds to thousands of different individual hybridization experiments can be performed simultaneously. This allows assays of enormous complexity to be carried out – for example, an analysis of the entire gene expression

profile of a cancer cell – with simplicity unimaginable only a few years ago. As a consequence many patents as well as scientific publications have accumulated during the last years. U.S. 6,194,158 discloses characteristic genes and gene expression useful in screening for, diagnosis of, monitoring 5 of, and therapeutic treatment of brain cancer. U.S. 6,218,122 discloses methods for determining or monitoring the progression of disease states or the efficacy of therapeutic regimens within human patients. U.S. 6'077'673 discloses mouse arrays having a plurality of probe polynucleotides corresponding to a key mouse gene for expression analysis of critical 10 mouse genes. A list of representative scientific papers dealing with monitoring the expression level of a large number of transcripts within a cell at any time are as follows: Schena et al., 1995, Quantitative monitoring of gene expression patterns with a complementary DNA-microarray, Science 270: 467-470; Lockhart et al., 1996, Expression monitoring by hybridization 15 to high-density oligonucleotide arrays, Nature Biotechnology 14:1675-1680; Blanchard et al., 1996, Sequence to array: Probing the genome's secrets, Nature Biotechnology 14:1649. Qi et al., 2003, Identification of genes responsible for osteoblast differentiation from human mesodermal progenitor cells PNAS 18;100(6):3305-10. While this list of scientific papers 20 and patents reflects without any doubt the great potential of microarrays, there are a couple of yet unsolved problems that are more and more discussed among the scientific community. Especially, these problems are data overflow, representative sample collection, RNA processing and inappropriate data analysis. It is even suspected that within next five years, 25 many of conclusions drawn from published data will be revised or refuted. Thus there remains a real and unmet need for advanced microarray solutions, targeted to specific tissues above all with respect to simplification and substantiation of the process of data generation and data handling. With respect to this issue the disclosed invention has made considerable 30 contribution in the cartilage area with a cartilage-specific microarray containing a manageable number of cartilage relevant genes.

Limitation on the number of cartilage relevant genes

Until today the number of cartilage-relevant genes (genes that have been associated a potential functional role on cartilage biology, homeostasis or pathology) is very limited. Approximately, 100-200 genes 5 have been described in the literature in any relationship to cartilage tissue. While existing publications e.g. Heller et al PNAS, 94; 2150-2155; 1997 have described analysis of inflammatory diseases of cartilage and Sekiya et al PNAS 99; 4397-4402; 2001 cartilage formation from stem cells with microarrays, a comprehensive analysis and determination of characteristic 10 gene expression profiles for 2D, 3D, fetal, adult and pathological chondrocytes cell cultures cultivated under different conditions has not been performed up to now. While in patent WO01/24833 A2 a few markers have been determined that are associated with chondrocytes and their phenotype stability, it will not be possible to perform a detailed gene 15 expression analysis and to define specific fingerprints. Therefore the possibility of characterizing culture conditions or cartilage tissue samples can not be thoroughly addressed.

Completion of the human genome first project draft on 2000 has revealed that the human genome comprises ~30000-35000 human 20 genes. Estimates show that the number and type of active genes vary significantly between different tissues and may increase up to a couple of 10000 for complex tissues, e.g. brain. As a consequence, many genes albeit fully sequenced may have yet not been disclosed to be functionally up- or down regulated in cartilage or cartilage derived cells. The inventive 25 approach described herein has made possible to up to now disclose a total of 467 known and additional genes being differentially expressed in a significant and objective manner within chondrocytes or chondrogenic cells.

By means of the already known and additionally found to be 30 cartilage related genes, a strategy to best address and represent chondrocytes cultured under different conditions has been developed in the scope of the present invention.

Summary of the Invention

In a first aspect the present invention relates to a method for the identification of tissue/cell specific marker genes comprising

- 5 a) taking tissue and/or cells of at least one developmental stage and/or at least one disease state, and/or
 cultivating said tissue and/or cells *in vitro* under at least one culture condition,
- 10 b) determination of gene expression profiles of said tissue/cells and/or *in vitro* cultivated tissue/cells and
- 15 c) identification of specific marker genes by bioinformatic analysis of said gene expression profiles.

In particular, the first aspect relates to a method for the identification of tissue/cell specific marker genes comprising cultivating tissue/cells of different developmental stages and/or health conditions *in vitro* under different culture conditions, determination of gene expression profiles of said *in vitro* cultivated cartilage tissue and identification of specific marker genes by bioinformatic analysis of said gene expression profiles.

In a preferred embodiment said tissue is selected from the group consisting of fetal tissue, adolescent tissue, adult tissue, healthy tissue, pathological tissue, progenitor cells such as stem cells or cells derived from the same precursor lineage. Preferred culture conditions are 2D and 3D *in vitro* cultures and the gene expression profiles are preferably determined by means of a micro-array. The bioinformatic analysis of said gene expression profiles is preferably done by cluster software such as e.g cluster analysis.

In a preferred embodiment said tissue is cartilage.

A second aspect of the present invention relates to a method for the determination of a disease state or developmental status of cells/tissue or the physiological potential of cells/tissue. Said method comprises establishing a profile of cellular constituents, preferably a gene expression profile, of said cells or tissue, comparison of said resulting gene expression profile with gene expression profiles characteristic for a particular status or physiological potential of the examined cells or tissue.

Said method can e.g. be used to assess the redifferentiation potential of cells or tissue, the assessment of the quality of tissue biopsies for diagnostic and prognostic purposes regarding *in vitro* tissue engineering applications, the assessment of the quality of *in vitro* produced cells such as 5 e.g. mesenchymal cells, stem cells or embryonic cells or of *in vitro* produced tissue for therapeutical applications and for determining the effect of one or more growth factors, media compositions or drugs on cells or tissue. Based on said method it is e.g. possible to set up different *in vitro* culture conditions for cells/tissue allowing the cultivation of cells/tissue which retain their 10 potential for differentiation.

In a preferred embodiment said cells or tissue is cartilage tissue or chondrocytes and the array comprises polynucleotide probes of tissue specific marker genes.

15 In a further preferred embodiment said profile is a gene expression profile which is determined by means of a micro-array.

A further object of the present invention is a method for the determination of characteristic profiles for clinical use comprising correlating the patient data of the biopsy donor with the gene expression profile of said 20 biopsy cells/tissue. Preferably said gene expression profile has been determined according to the above disclosed method. The resulting profiles of said method are suitable tools in the clinic allowing an evaluation of further treatments of a patient.

The present invention provides characteristic gene expression profiles experimentally determined by using cartilaginous 25 tissues as from individual human donors of various ages (fetal, adolescent, adult) and health conditions (healthy and arthritic) or cells thereof cultivated under different *in vitro* culture conditions (2D and 3D *in vitro* cultures, time follow ups). From these different gene expression profiles a set of hitherto 467 markers has been deduced that can be used to design and produce a 30 cartilage specific microarray for commercial applications in the field of R&D, such as culture media development, drug screening etc., but also for clinical applications.

Gene expression analysis performed with such microarrays and the corresponding analytical procedure thereof can be used to assess quality control of human donor cartilage, e.g. biopsy and therefore optimization of any downstream tissue engineering process, for diagnostic evaluation of the patient and its candidate treatment methods, to ensure a cost-optimized procedure, to investigate and assess all kind of 2D- and 3D *in vitro* cultures performed with human chondrocytes or chondrogenic cells, e.g. stem cells, to screen all kind of drugs, e.g. hormones, growth factors within the above mentioned *in vitro* cultures regarding a potential beneficial effect and quality assessment of *in vitro* produced tissue performed by tissue engineered procedures.

In a further aspect the present invention provides a cartilage array comprising a plurality of different polynucleotide probe spots stably associated with a solid surface of a carrier, whereby each of said spots is made of a unique polynucleotide that corresponds to one specific cartilage marker gene.

A preferred cartilage array of the present invention comprises at least two spots that have different nucleotide sequences but of the same cartilage marker gene, more preferably at least 10 spots indicative for one tissue or cell status, whereby said at least 10 spots can be selected from different sequences of one gene or from different genes or a combination thereof.

In a preferred embodiment said polynucleotides of the array do not cross hybridize under stringent conditions with each other.

In a preferred embodiment of the present invention the cartilage array comprises spots that are indicative for at least two tissue or cell status, preferably 3.

A further preferred inventive cartilage array is an array wherein at least part of the cartilage marker genes are selected from the 467 genes listed in the description, preferably at least 10 %, more preferably at least 50 %, most preferably about 100 %.

A further preferred inventive cartilage array is an array wherein at least part of the cartilage marker genes are selected from a

subgroup of the 467 genes listed in the description, wherein said subgroup consists of the most tissue specific 200 genes.

In another preferred embodiment the status is selected from biopsies and/or 2D cultures and/or 3D cultures of healthy adult, healthy fetal/infant, undesired adult, undesired fetal/infant or progenitor cells like e.g. stem cells or cells derived from the same precursor lineage.

In a further preferred embodiment of the present invention the polynucleotide probes of the cartilage array have a length of at least 10 nucleotides, preferably at least 20 nucleotides. The probes can also have a length of 30 nucleotides, 50 nucleotides or 70 nucleotides. It is as well possible to use PCR derived products produced from cDNA clones.

In a preferred embodiment the carrier of the inventive cartilage array is attached to coated glass, nylon or any other material.

A further object of the present invention is a kit for use in a hybridization assay, wherein said kit comprises a cartilage array of the present invention. In a preferred embodiment said kit comprises reagents for generating a labelled target polynucleotide sample, a hybridization buffer and a wash medium.

20 Description of the Figures

The present invention will be further understood from the following description with reference to the tables and figures where:

Tab.I shows the determined number of all genes in the corresponding SOM analysis being differentially expressed according to microarray analyses of a variety of *in vitro* chondrocyte cultures according to predefined criteria. From these data sets specific expression profiles can be deduced that are characteristic for different cell culture conditions.

Tab.II shows the extracted and reviewed genes deduced from Tab I in order to have only single entry numbers. Since most of these genes have never been described in any relationship to cartilage, they can be considered as novel cartilage marker (positive/negative markers) or key cartilage genes.

Tab III shows a subset of marker genes from Tab. II that has been used for the production of a micro-array. Included is a subset from Tab II and genes known from the literature.

5 Tab IV shows the results of the analysis of the 467 cartilage specific marker genes.

Tab V shows the samples used in Examples 1, 2 and 3. Human chondrocytes isolated from 4 different donors were proliferated and kept in 3D-like pellet culture for 7 and 14 days resulting in a total number of 12 samples.

10 Fig.1 shows a classical result from an analysis performed with self-organizing-maps. This software clusters all genes together in sub clusters that show a similar expression profile. The number of marker genes for the corresponding analysis e.g. 2D vs. 3D cultures (see also Tab I) corresponds to the total number of genes in the sub clusters.

15 Fig 2 shows an example of a graphical presentation of a cluster analysis and viewed by the software treeview. This shows how cells from different origin and potential for *in vitro* cartilage formation are related to each other and allow a clearer classification of the cell sources. Fetal cells clearly produce different gene clusters compared to adult chondrocytes,
20 while failures are characterized by other gene clusters. Furthermore 3D cell cultures analyzed in a time dependent manner from different donors can be distinguished among each other and gene expression profiles will be grouped accordingly.

25 Fig 3: SOM analysis of all culture conditions and samples described in Example 2 and in Tab V.

Fig 4: SOM analysis for proliferated chondrocytes (t0) only, for the 4 donors. Gene expression pattern corresponding to donor 2 (the second spot from left hand side in every cluster) behaves different in most clusters.

30 Fig 5: SOM analysis of chondrocytes kept in 3D culture condition for 7 days (t7). Gene expression pattern from donor 3 (the third spot from left hand side in every cluster) is different for example in clusters c2 and c5.

Fig 6 shows self organized maps (SOM) of chondrocytes from same patients of Figures 4 and 5 kept under 3D culture condition for 14 days (t14).

Fig 7: cluster analysis of all culture conditions and samples described in Example 2 and in Tab V. This figure shows a subset of 88 hierarchical clustered genes (rows) and samples (columns) demonstrating similar gene expression behavior of chondrocytes under different culture conditions. For example proliferated cells (#1, #2, #4, #5, #7, #8, #10, #11) can easily be discriminated from cells kept in 3D-like pellet culture for 14 days (3#, 6#, 9#, 12#).

Fig 8: cluster analysis of human aortic fibroblasts vs. chondrocytes. This figure shows a subset of selected clusters of human aortic fibroblasts cells compared to human chondrocytes both kept in 3D pellet cultures for 14 days. The dendrogram in the upper part of the figure shows the ability of CART-CHIP™ 300 microarray described in this invention to discriminate between different cell lines.

Fig 9: cluster analysis of Interleukin-1 treated vs. untreated human chondrocytes. This figure demonstrates a subset of representative gene clusters allowing differentiation between cells treated with Interleukin-1 from untreated cells both kept in 3D pellet cultures as well as for proliferated cells.

Detailed Description of the Invention

Definitions

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2D cultures as used in the scope of the present invention are anchorage dependent chondrocyte cultures cultivated on plastic culture devices.

30 *3D cultures* as used in the scope of the present invention are chondrocytes cultured in a three dimensional environment, namely either a) scaffold-free, such as small high density pellet cultures ($0.25\text{-}3.0 \times 10^6$ cells) or as high density cultures using 50×10^6 cells/ml or aliquots thereof; or b) by using a synthetic scaffold such as PGA, PLA, or mixtures

thereof or biological substances such as agarose, alginate, chitosan or collagen.

failures as used in the scope of the present invention are chondrocytes cultured in a three dimensional environment that are not able 5 to synthesize new extracellular matrix thereby compromising the production of new living tissue engineered cartilage equivalents.

gene expression profile as used in the scope of the present invention is a profile of genes that are up or down regulated according to different cell conditions.

10 *fingerprint* as used in the scope of the present invention refers to a gene expression profile characteristic for a cellular status.

tissue or cell status as used in the scope of the present invention refers to a tissue or cells therof having a certain metabolic or activity status.

15 *new extracellular matrix* as used in the scope of the present invention designates living cartilage-like tissue.

micro-array as used in the scope of the present invention is used in its original scope that encompasses embodiments today sometimes referred to as "macro-arrays".

20 The Invention

The present invention provides cartilage-specific gene arrays as well as methods for their use. In the subject cartilage arrays, a plurality of polynucleotide probe spots are stably associated with the 25 surface of a solid carrier, preferably a surface of a microscope glass slide. Each different polynucleotide probe spot is made of a unique polynucleotide that corresponds to a key cartilage gene of interest. Thus, the subject arrays find particular use in gene expression assays of key cartilage genes. In further describing the subject of the invention, the cartilage specific 30 microarrays are first discussed, followed by a review of representative applications in which the subject arrays may be employed.

Arrays of the Subject Invention-General Description**Selection of novel key cartilage specific genes:**

A critical feature of the subject arrays is that all of the probe polynucleotide spots of the array correspond to human key cartilage genes
5 that have been found through unique selection processes and criteria. As a result of said processes, up to now 467 different key human cartilage genes that are under tight transcriptional role have been discovered, some of them being not described before in any relationship to cartilage. In more detail, different microarray analyses were performed by using cartilaginous tissues
10 as from individual human donors of various ages (fetal, adolescent, adult) and health conditions (healthy and arthritic) or cells thereof cultivated under different *in vitro* culture conditions (2D and 3D *in vitro* cultures, time follow ups). This variety of cartilage cell sources and different culture conditions was set up to grasp the highest possible number of genes differentially
15 expressed and thus being indicative of a potential role.

It has been found that specific chondrocyte culture conditions are of great importance for the present invention that discloses a plurality of novel key cartilage genes as well as characteristic and meaningful gene expression patterns. For this reason, the strategy and
20 criteria of the analysed *in vitro* human chondrocyte cultures are described in more detail. The principal experimental setup included both the cultivation of chondrocytes in an anchorage dependent condition, known as 2D cultures for expansion of cells e.g. where the passages is variable but at least more than one, as well as cultivation of chondrocytes in an anchorage
25 independent condition, known as 3D cultures for (re-)differentiation and de novo tissue formation of cells. These are the key steps of any tissue engineering process where autologous tissue equivalents are produced. Since the cell source is either a small biopsy, a small bone marrow aspirate in case of mesenchymal stem cells or other tissue with a limited number of
30 pre-chondrogenic cells, it is first necessary to isolate those cells in order to be able to multiply the cell number drastically. In case of a cartilage biopsy, cells are released from their surrounding extracellular matrix by collagenase digestion and then seeded onto the surface of plastic tissue culture flasks.

The proliferation may take place either in the presence or absence of fetal serum combined with conventional DMEM/F12 medium. Cells can then be passaged by trypsin treatment over several rounds. As a major drawback of this necessary cell expansion, the cells loose their differentiated phenotype 5 and assume a de-differentiated phenotype with altered gene expression. It is further known that with increasing number of passages the state of de-differentiation also advances. As a consequence, genes being transcriptionally upregulated under such artificial culture conditions are cartilage relevant in a manner being indicative of an undesired cellular 10 status. It is also quite common to designate these genes as de-differentiation or negative markers. While healthy tissue in general has been found to re-differentiate in 3D culture after up to 4 passages in 2D cultures, tissue of undesired cellular status cultivated under usual conditions, such as usual culture media, usually does not re-differentiate in 15 3D culture after at most 4 passages in 2D culture.

Subculture modulated chondrocytes that do not express differentiation markers reexpress the differentiated phenotype in response to the anchorage- independence resulting from various 3D culture models, e.g. high density cultures, agarose or alginate cultures, or cultures within 20 synthetic scaffolds such as made of polyglycolic acid (PGA), polylactic acid (PLA) or mixtures thereof. To set up three dimensional cell cultures the cells are detached after proliferation by trypsin treatment and embedded either in gel-like substances such as alginate, seeded within a porous scaffold such as PGA or cultivated as high-density cultures, only. The time 25 for the analysis may vary and ideally addresses several time points (up to several weeks). Thus 3D *in vitro* chondrocyte cultures support the differentiated phenotype of chondrocytes and can be used to discover cartilage relevant genes or differentiation markers. It should be noted however, that reversibility of the de-differentiation process is dependent on 30 the number of passages and can become irreversible or at least partially irreversible at higher passage numbers (under usual conditions at most about 4 passages). As a rule the time course of de- and re-differentiation are similar. During skeletal development, cartilage serves as a template for

bone formation. Chondrocytes of fetal or infant (< 1 year) or growth plate cartilage pass through different stages and exhibit several distinct phenotypes, such as resting, proliferating, and hypertrophic chondrocytes. Progression through each of these phases is accompanied by profound changes in gene expression patterns. Further, evidence has accumulated that the successful sequence of cartilage repair via tissue engineering recapitulates aspects of embryonic tissue formation. For these reasons, it is important to consider fetal and infant cartilaginous tissue. Cells isolated from human fetal/infant cartilage that are cultivated in 2D and 3D culture systems as described above are especially helpful to understand the mechanisms underlying the phenotypic instability of chondrocytes and the related gene expression patterns. These 2D and 3D culture system may then be analyzed to deduce gene expression profiles and to define marker genes that are characteristic for the (re-)differentiation process. Thus maintenance of chondrocyte-specific phenotype being crucial for normal structure and biomechanical properties of articular cartilage may be better understood and have important implications for modern therapeutic biological applications.

The above mentioned experimental setup for 2D and 3D cultures may be even expanded to compare human adult cells with human fetal/infant chondrocytic cells of age <1 year. The comparison of gene expression profiles of adult versus fetal/infant human chondrocytes during the *in vitro* cartilage formation process is an important aspect since marker genes associated with developmental aspects are revealed. This can be of further interest when 3D cell cultures need to be optimized for their *in vitro* performance for the production of new tissue by e.g. adding growth factors that are found to play a major role during the early onset of cartilage formation *in vivo*.

Another experimental setup found in the scope of this invention includes the *in vitro* culture of cells harvested from cartilaginous areas of arthritic knee joints. Osteoarthritis (OA) results from the failure of chondrocytes within the joint to maintain the balance between synthesis and degradation of extracellular matrix. OA is a multifactorial disorder in

which aging, genetic, hormonal and mechanical factors are all major contributors to its onset and progression. With progressing disease state, the articular chondrocytes ability to maintain homeostasis and functionality is increasingly disappearing. As a consequence, the phenotype of 5 osteoarthritic chondrocytes compared with normal chondrocytes exhibits remarkable changes. Gene expression profiling allows characterization of the osteoarthritic cellular phenotype, a key determinant for understanding and manipulation of osteoarthritic processes. By studying and comparing the gene expression profiles of chondrocytes harvested from pathological 10 and healthy human cartilage areas it becomes possible to identify marker genes that are able to predict the future outcome of cell cultures used for *in vitro* tissue engineering applications. This also relates to the very critical question of the assessment of the quality of the starting biopsy material that is being used for downstream applications like tissue engineering. By 15 having this important information before performing any downstream applications like e.g. proliferation and consecutive 3D *in vitro* tissue formation, the further steps of any process can then be adapted or even not performed at all because of inadequate quality of biopsy material. Such decision may be of high relevance when tissue-engineering processes are 20 transferred or applied in the clinic. Gene expression profiling of chondrocytes may then be used as a diagnostic tool to allow and to choose that therapeutic approach with the most promising clinical outcome.

A further important aspect of the invention is the observation that chondrocytes derived from osteoarthritic patient material 25 always qualify for anchorage dependent proliferation in 2D over several passages. These cells however, if subsequently induced to re-differentiate by culturing them as 3D high density pellets, do not survive over an extended time period, in most cases they die in culture by undergoing apoptosis. It is assumed that these cells, due to an altered phenotype, are 30 not capable of producing the critical survival factors in the appropriate concentrations, above all extracellular matrix components providing intercellular spaces as they occur in native cartilage. Cells that are not suitable to be cultured within 3D high density cultures are herein referred to

as „failures“. These impaired cell cultures can be used to set up representative “failure” systems, where cells from different pathological cartilage sources are harvested, proliferated and cultivated in 3D high density pellet culture systems. After each of these experimental steps, RNA 5 can be isolated from the different cell sources and combined to create “failure pools”. These failure pools are very well suitable to identify general marker genes being indicative of the onset of osteoarthritis.

For finding cartilage relevant genes, and for determining their presence dependent on the specific cartilage type such as age, health 10 etc., sufficient material must be generated, e.g. by 2D culturing over several passages, and optionally 3D culturing. Said material then can on be subjected to usual gene analyses, and the tissue specific genes determined. Cartilage samples are classified prior to culturing and/or after culturing to get the information needed for later interpretation of the gene 15 expression profile.

A further experimental setup of the current inventions discloses the analysis of chondrocytes grown in 3D cultures isolated from pathological human cartilage and analyzed in a time dependent manner. This experimental set-up allows to study the apoptotic process and to 20 further define additional dynamic and characteristic gene expression profiles, useful for deducing and further assessment of the quality of the biopsy material.

The microarray process and strategy for disclosing all the cartilage relevant genes with the above-mentioned tissues and cell culture 25 criteria will be described in the following. An important issue of the innovative strategy used by the inventors of the present inventions is to use various microarrays containing a high number of genes comprising different functional categories preferentially by representing the whole genome. The broader the microarray regarding the coverage of the human genome the 30 more genes associated with chondrocyte cell cultures can be determined. The chosen strategy of the inventors was not obvious to a person skilled in the art.

RNA isolated from the above mentioned different cell cultures conditions may be radioactive labeled with e.g. 33P or fluorescence like e.g. Cy3 and hybridized to the corresponding filters or microarrays. After hybridization each array may then be scanned and the 5 corresponding signals measured (Tab IV). This raw data file needs then to be calibrated and normalized in a manner to create an input file for the further downstream analysis process. In principle if the data are normalized an expression profile is created. To identify the key cartilage marker genes being differentially expressed under the chosen criteria, tedious 10 bioinformatic analysis are conducted. Corresponding cell cultures and their expression profiles are therefore compared and analyzed accordingly and the different clusters of marker genes determined by software analysis e.g. self-organizing maps (herein referred to as SOM). A representative example of a result for the comparison of different gene expression profiles 15 from different cell culture conditions performed by SOM analysis is given in Fig.1. By performing SOM analysis genes that are similarly expressed are clustered together in so-called sub clusters. The total amount of marker genes for one analysis corresponds to the total amount of sub clusters containing the corresponding genes. Table I in the appendix summarize the 20 results of all the different analysis performed and encompasses all the genes determined for every set of cell culture analysis.

By performing this analytical procedure the analysis reveals several characteristic up and down regulated marker genes for different cellular culture conditions. From these marker genes characteristic 25 expression profiles can then be deduced and used as a benchmark for the comparison or further characterization of other cell cultures.

Hence, on the one hand previously unknown cartilage-relevant genes associated with different culture conditions and on the other hand characteristic gene expression profiles (cellular fingerprints) indicative 30 of a stage of development, a disease state or a particular selected cell culture condition are revealed. These fingerprints are part of the current invention and are of major importance for the classification and

characterization of chondrocytes cultivated under different culture conditions.

Since the gene clusters from all the different analysis contain repetitive gene entries, they have been further processed so that
5 only single entry genes are recorded (see Tab II). This 467 selected sequences are thus all key cartilage genes that are activated and thus differentially expressed according to a stage of development, a disease state or a particular selected cell culture condition and are part of the current invention. A list of all 467 genes with their Pubmed accession no.
10 and a description is given below See also Tables II and III):

List of Table II related sequences:

Pubmed Accesion No	Description
AA283693	Human osteoclast stimulating factor mRNA, complete cds
AA845156	Serine protease inhibitor, Kazal type 1
R52548	Human superoxide dismutase (SOD-1) mRNA, complete cds
T67128	ARYLAMINE N-ACETYLTRANSFERASE, MONOMORPHIC
AA845015	Elastase 1, pancreatic (elastase IIA)
AA937895	Antigen identified by monoclonal antibodies 12E7, F21 and O13
AA844998	Pancreatic polypeptide
AA844818	Amylase, alpha 2A; pancreatic
AA894557	Creatine kinase B
AA872001	Annexin VI (p68)
H09590	Human mRNA for eukaryotic initiation factor 4AI
AA868278	Testis specific protein 1 (probe H4-1 p3-1)
AA490855	Acid finger protein ZNF173
H05820	Human MRL3 mRNA for ribosomal protein L3 homologue (MRL3 = mammalian ribosome L3)
N57766	Agammaglobulinaemia protein-tyrosine kinase atk
AA873885	Alkaline phosphatase, liver/bone/kidney
AA878880	Interferon (gamma)-induced cell line; protein 10 from
R54818	Human eukaryotic initiation factor 2B-epsilon mRNA, partial cds
AA458630	RENIN PRECURSOR, RENAL
W37864	Phosphatase and tensin homolog (mutated in multiple advanced cancers 1)
N63192	Phenylethanolamine N-methyltransferase
R55789	Human X11 protein mRNA, partial cds
R56871	Human chromatin assembly factor-I p60 subunit mRNA, complete cds
AA448659	M-PHASE INDUCER PHOSPHATASE 2
AA235388	Tropomodulin
W37769	Chromogranin B (secretogranin 1)
AA421701	H.sapiens mRNA for MUF1 protein
N81029	Collagen, type XVIII, alpha 1
AA644128	Nuclear autoantigenic sperm protein (histone-binding)
N26536	ATPase, Cu++ transporting, beta polypeptide (Wilson disease)
AA890663	Human protein kinase PAK1 mRNA, complete cds
AA405987	Glycerol kinase 2 (testis specific)
AA888182	Ribosomal protein S4, X-linked
H09730	Adenylate kinase 2 (adk2)
AA285155	CDC46 HOMOLOG
AA873351	Ribosomal protein L35a
H12320	CAMP-RESPONSE ELEMENT BINDING PROTEIN
AA856556	Ribosomal protein S28
R43581	Human guanine nucleotide-binding protein G-s, alpha subunit mRNA, partial cds
AA633768	60S RIBOSOMAL PROTEIN L24
AA496880	Ribosomal protein L5
AA625632	Ubiquitin A-52 residue ribosomal protein fusion product 1
R40850	H.sapiens mRNA for alpha-centractin
AA486072	Small inducible cytokine A5 (RANTES)
N80129	Metallothionein 1L
T67270	UBIQUINOL-CYTOCHROME C REDUCTASE COMPLEX SUBUNIT VI REQUIRING PROTEIN
AA775364	60S RIBOSOMAL PROTEIN L30
AA464743	Ribosomal protein L21
AA663983	Triosephosphate isomerase 1
AA634008	40S RIBOSOMAL PROTEIN S23

AA683050 40S RIBOSOMAL PROTEIN S8
 AA775874 60S RIBOSOMAL PROTEIN L18
 AA029934 Integrin, alpha V (vitronectin receptor, alpha polypeptide, antigen CD51)
 AA872397 GALECTIN-2
 AA428195 Protein tyrosine phosphatase, non-receptor type 2
 AA478724 Insulin-like growth factor binding protein 6
 T40541 H.sapiens mRNA for human giant larvae homolog
 N33214 H.sapiens mRNA for membrane-type matrix metalloproteinase 1
 W69399 Homo sapiens adenosine triphosphatase mRNA, complete cds
 H85454 Homo sapiens delayed-rectifier K⁺ channel alpha subunit (KCNS1) mRNA, complete
 cds
 T71284 Complement component 1, q subcomponent, beta polypeptide
 N95418 Human FK-506 binding protein homologue (FKBP38) mRNA, complete cds
 AA430675 Human DNA repair protein XRCC9 (XRCC9) mRNA, complete cds
 AA682851 Homo sapiens mRNA for ERp28 protein
 AA427433 PROTEIN PHOSPHATASE PP2A, 65 KD REGULATORY SUBUNIT, ALPHA
 ISOFORM
 AA100296 H.sapiens PAP mRNA
 AA070997 Proteasome (prosome, macropain) subunit, beta type, 6
 R27585 Proteasome component C2
 N71628 Spi-B transcription factor (Spi-1/PU.1 related)
 AA464566 Human mRNA for LDL-receptor related protein
 AA043228 Calponin 3, acidic
 AA478273 APEX nuclease (multifunctional DNA repair enzyme)
 H05619 Homo sapiens GDNF family receptor alpha 2 (GFRalpha2) mRNA, complete cds
 AA405562 Protein phosphatase 4 (formerly X), catalytic subunit
 AA147043 Homo sapiens CAGH1a (CAGH1) mRNA, partial cds
 AA035384 Homo sapiens mRNA for small subunit of cytochrome b in succinate dehydrogenase
 complex, complete cds
 R60150 Human mRNA for histidyl-tRNA synthetase (HRS)
 N64051 Homo sapiens Werner syndrome gene, complete cds
 AA405748 SPLICING FACTOR U2AF 65 KD SUBUNIT
 AA461110 Homo sapiens growth-arrest-specific protein (gas) mRNA, complete cds
 AA845167 ELASTASE IIIA PRECURSOR
 AA443118 Homo sapiens mRNA for CD151, complete cds
 N92319 Glycoprotein Ib (platelet), beta polypeptide
 AA187148 Core-binding factor, beta subunit
 AA253413 Friedreich ataxia
 AA046701 ATP SYNTHASE LIPID-BINDING PROTEIN P1 PRECURSOR
 AA164562 Homo sapiens actin-related protein Arp3 (ARP3) mRNA, complete cds
 AA496357 Homo sapiens SKB1Hs mRNA, complete cds
 AA180742 TUBULIN ALPHA-4 CHAIN
 AA454743 Human protease M mRNA, complete cds
 AA437226 Interleukin 10 receptor
 AA458849 Homo sapiens placental bikunin mRNA, complete cds
 AA504891 Crystallin, alpha B
 AA609655 Homo sapiens mRNA for SCP-1, complete cds
 AA599158 MULTIFUNCTIONAL AMINOACYL-TRNA SYNTHETASE
 AA052932 Homo sapiens casein kinase I gamma 2 mRNA, complete cds
 AA789328 Homo Sapiens (clone PK2J) CDC2-related protein kinase (PISSLRE) mRNA,
 complete cds
 AA129537 Human GAP SH3 binding protein mRNA, complete cds
 AA486209 Low density lipoprotein-related protein-associated protein 1 (alpha-2-macroglobulin
 receptor-associated protein 1
 H39018 H.sapiens Syt V gene (genomic and cDNA sequence)
 AA464217 V-akt murine thymoma viral oncogene homolog 1

T95053 Homo sapiens Rigui (RIGUI) mRNA, complete cds
 AA454646 LYMPHOTOXIN-BETA RECEPTOR PRECURSOR
 AA448400 Human plectin (PLEC1) mRNA, complete cds
 H13691 Major histocompatibility complex, class II, DM beta
 AA132086 Homo sapiens RCL (Rcl) mRNA, complete cds
 AA488073 Mucin 1, transmembrane
 N40945 H.sapiens mRNA for DRES9 protein
 R55705 Homo sapiens orexin receptor-1 mRNA, complete cds
 H50114 Homo sapiens NMDA receptor mRNA, complete cds
 AA452841 Human K-Cl cotransporter (hKCC1) mRNA, complete cds
 W73790 IMMUNOGLOBULIN-RELATED 14.1 PROTEIN PRECURSOR
 N30302 POSSIBLE GTP-BINDING PROTEIN HSR1
 AA291556 Human ras inhibitor mRNA, 3' end
 AA598510 Human APRT gene for adenine phosphoribosyltransferase
 AA453787 Human TFIIB related factor hBRF (HBRF) mRNA, complete cds
 H05655 Human transcriptional activator mRNA, complete cds
 AA419177 INTEGRAL MEMBRANE PROTEIN E16
 AA458807 Human retinal protein (HRG4) mRNA, complete cds
 AA293218 Cleavage stimulation factor, 3' pre-RNA, subunit 2, 64kD
 W44860 Human calmodulin mRNA, complete cds
 AA629862 Homo sapiens mRNA for smallest subunit of ubiquinol-cytochrome c reductase, complete cds
 AA447674 Homo sapiens HIV-Nef associated acyl CoA thioesterase (hNAACTE) mRNA, complete cds
 T52484 Nerve growth factor beta
 AA496810 Protein kinase C substrate 80K-H
 AA486233 G1 to S phase transition 1
 AA079775 TYROSINE-PROTEIN KINASE CSK
 W73889 Tetranectin (plasminogen-binding protein)
 R50337 Solute carrier family 19 (folate transporter), member 1
 R55046 MpV17 transgene, murine homolog, glomerulosclerosis
 R46821 T-COMPLEX PROTEIN 1, ALPHA SUBUNIT
 R87763 Human telencephalin precursor mRNA, complete cds
 H69583 Human BTG2 (BTG2) mRNA, complete cds
 R56046 Guanine nucleotide binding protein (G protein), alpha z polypeptide
 AA922705 Glycogen phosphorylase B (brain form)
 AA487571 Surfactant, pulmonary-associated protein C
 AA402440 Homo sapiens exportin t mRNA, complete cds
 H29521 ATP-binding cassette 3
 AA490911 Homo sapiens drp1 mRNA, complete cds
 AA486082 Homo sapiens sgk gene
 AA678065 2,3-bisphosphoglycerate mutase
 R43509 Human Gu binding protein mRNA, partial cds
 N57553 Adenosine receptor A2
 AA676955 Aplysia ras-related homolog 12
 R14692 Human Na/H antiporter (APNH1) mRNA, complete cds
 AA488979 Homo sapiens nucleolar protein (MSP58) mRNA, complete cds
 AA443630 Aldehyde dehydrogenase 8
 AA027840 H.sapiens mRNA for RIT protein
 AA456830 Diacylglycerol kinase, alpha (80kD)
 AA453015 H.sapiens L23-related mRNA
 AA074446 Human GTP cyclohydrolase I feedback regulatory protein gene, complete cds
 AA027042 DNA-DIRECTED RNA POLYMERASE II 23 KD POLYPEPTIDE
 AA629923 Human mRNA for pM5 protein
 AA460830 Homo sapiens (clone mf.18) RNA polymerase II mRNA, complete cds

AA454218 Homo sapiens transcription factor SL1 mRNA, complete cds
 AA046523 H.sapiens mRNA for centrin gene
 R51346 Human eIF-2-associated p67 homolog mRNA, complete cds
 AA029964 Human ataxin-2 related protein mRNA, partial cds
 AA489219 DUTP pyrophosphatase
 AA043133 Solute carrier family 16 (monocarboxylic acid transporters), member 1
 AA812973 Human mRNA for testis-specific TCP20, complete cds
 AA453471 GANGLIOSIDE GM2 ACTIVATOR PRECURSOR
 AA284693 Transcription factor AP-4 (activating enhancer-binding protein 4)
 N90281 Human B7 mRNA, complete cds
 AA629542 Brush-1
 AA679345 Human BTK region clone ftp-3 mRNA
 H37774 Tuberin
 T97181 Platelet factor 4
 AA454879 Plasminogen activator, urokinase receptor
 AA147640 Phosphorylase, glycogen; liver (Hers disease, glycogen storage disease type VI)
 AA757429 Human serotonin N-acetyltransferase mRNA, complete cds
 AA490991 Homo sapiens HnRNP F protein mRNA, complete cds
 AA422058 H.sapiens mRNA for D1075-like gene
 N66208 Human (ard-1) mRNA, complete cds
 AA630776 Human AP-3 complex delta subunit mRNA, complete cds
 AA827287 Human interferon-induced leucine zipper protein (IFP35) mRNA, partial cds
 AA488084 Superoxide dismutase 2, mitochondrial
 R89715 Protein kinase C, gamma
 AA490501 H.sapiens mRNA; UV Radiation Resistance Associated Gene
 N32199 Human melanoma antigen recognized by T-cells (MART-1) mRNA
 AA434404 DNA primase polypeptide 2A (58kD)
 N93686 Aldehyde dehydrogenase 7
 AA292676 Human metargidin precursor mRNA, complete cds
 AA464417 INTERFERON-INDUCIBLE PROTEIN 1-8U
 AA442092 Catenin (cadherin-associated protein), beta 1 (88kD)
 AA026644 Transcription factor 3 (E2A immunoglobulin enhancer binding factors E12/E47)
 AA481464 Peptidylprolyl isomerase B (cyclophilin B)
 T68859 Alpha-2-plasmin inhibitor (alpha-2-PI)
 AA699560 Surfeit 1
 AA705069 Human mRNA for receptor of retinoic acid
 AA457739 Homo sapiens putative OSP like protein mRNA, partial cds
 H99843 Homo sapiens mRNA for quinolinate phosphoribosyl transferase, complete cds
 AA399410 Signal transducer and activator of transcription 3 (acute-phase response factor)
 AA443039 HEAT SHOCK 70 KD PROTEIN 1
 AA164440 Human autoantigen pericentriol material 1 (PCM-1) mRNA, complete cds
 AA446453 Human mRNA for c-myc binding protein, complete cds
 AA280692 Diacylglycerol kinase delta
 AA031514 Matrix metalloproteinase 7 (matrilysin, uterine)
 R33154 Msh (Drosophila) homeo box homolog 1 (formerly homeo box 7)
 AA487452 Human DNA fragmentation factor-45 mRNA, complete cds
 AA400329 Human gene for neurofilament subunit M (NF-M)
 AA454668 Prostaglandin-endoperoxide synthase 1 (prostaglandin G/H synthase and cyclooxygenase)
 AA486393 Cytokine receptor family II, member 4
 R52541 unknown EST
 AA171613 Homo sapiens carbonic anhydrase precursor (CA 12) mRNA, complete cds
 AA235706 Human TATA-binding protein associated factor 30 kDa subunit (tafII30) mRNA, complete cds
 AA668527 Human mucosal addressin cell adhesion molecule-1 (MAdCAM-1) mRNA, complete

	cds
T54144	Homo sapiens homolog of the <i>Aspergillus nidulans</i> <i>sudD</i> gene product mRNA, complete cds
R14080	Calcium modulating ligand
AA609599	Homo sapiens SSX3 (SSX3) mRNA, complete cds
AA489201	H.sapiens mRNA for PHAPI2b protein
R08876	Human 26S proteasome-associated <i>pad1</i> homolog (POH1) mRNA, complete cds
H46425	H.sapiens Pur (pur-alpha) mRNA, complete cds
R56149	Human putative transmembrane protein (nma) mRNA, complete cds
AA454619	Homo sapiens mRNA for Hic-5, partial cds
H15445	H.sapiens mRNA for SEX gene
AA705225	Myosin, light polypeptide 4, alkali; atrial, embryonic
AA191488	Human high-affinity copper uptake protein (hCTR1) mRNA, complete cds
N64862	Human SLP-76 associated protein mRNA, complete cds
R45413	Human transmembrane 4 superfamily protein (SAS) mRNA, complete cds
R77293	Intercellular adhesion molecule 1 (CD54), human rhinovirus receptor
AA436187	Integrin, alpha M (complement component receptor 3, alpha; also known as CD11b (p170), macrophage antigen alpha polypeptide)
AA676470	H.sapiens IAI.3B mRNA
AA443634	Homo sapiens ubiquitin conjugating enzyme G2 (UBE2G2) mRNA, complete cds
AA664180	Glutathione peroxidase 3 (plasma)
W58658	H.sapiens mRNA for CLPP
H54023	Homo sapiens monocyte/macrophage Ig-related receptor MIR-10 (MIR cl-10) mRNA, complete cds
H73724	Cyclin-dependent kinase 6
T70031	Human neutral amino acid transporter B mRNA, complete cds
AA481758	DNAJ PROTEIN HOMOLOG 1
AA521431	Human profilin mRNA, complete cds
AA446103	ERGIC-53 PROTEIN PRECURSOR
N92646	Immunoglobulin gamma 3 (Gm marker)
AA453789	Protein-tyrosine kinase 7
AA425299	Homo sapiens ezrin-radixin-moesin binding phosphoprotein-50 mRNA, complete cds
AA868929	Troponin T1, skeletal, slow
R60019	Homolog 2 of <i>Drosophila</i> large discs
AA857343	Human putative RNA binding protein (RBP56) mRNA, complete cds
AA481438	Complement component 1 inhibitor (angioedema, hereditary)
AA399674	Human small proline rich protein (sprlI) mRNA, clone 1292
T98887	Glucose-6-phosphatase
AA676404	Peptidylprolyl isomerase C (cyclophilin C)
H15747	Human HU-K4 mRNA, complete cds
H16958	Human glyceraldehyde 3-phosphate dehydrogenase mRNA
AA936783	Eukaryotic translation initiation factor 3 (eIF-3) p36 subunit
AA884709	Cytochrome P450 11 beta
H24688	Human SWI/SNF complex 170 KDa subunit (BAF170) mRNA, complete cds
AA884403	Human cardiotrophin-1 (CTF1) mRNA, complete cds
AA404619	5' nucleotidase (CD73)
AA598611	IMMEDIATE-EARLY RESPONSE PROTEIN NOT
H72875	GATA-binding protein 3
H63361	Eukaryotic translation initiation factor 2B (eIF-2B) alpha subunit
R39221	Human MAP kinase mRNA, complete cds
R02346	U1 snRNP 70K protein
R51835	unknown EST
R33031	H.sapiens mRNA for sigma 3B protein
AA412053	CD9 antigen
AA001897	Erythroid alpha-spectrin

W81191	Homo sapiens nucleolar autoantigen No55 mRNA, complete cds
AA430552	Homo sapiens proline-rich Gla protein 2 (PRGP2) mRNA, complete cds
AA394130	Human transducin-like protein mRNA, complete cds
N92864	Human cleavage and polyadenylation specificity factor mRNA, complete cds
AA457123	VALYL-TRNA SYNTHETASE
R43320	Human guanine nucleotide-binding regulatory protein (Go-alpha) gene
AA670430	Glutamate receptor, metabotropic 3
H65066	Visinin-like 1
AA458785	GUANYLATE CYCLASE SOLUBLE, BETA-1 CHAIN
AA485871	H.sapiens mRNA for myosin-I beta
T39411	Human 53K isoform of Type II phosphatidylinositol-4-phosphate 5-kinase (PIPK) mRNA, complete cds
R00855	Homo sapiens 59 protein mRNA, 3' end
H98666	Metallopeptidase 1 (33 kD)
H72028	GELSOLIN PRECURSOR, PLASMA
AA679177	Human follistatin-related protein precursor mRNA, complete cds
N21576	Human mitochondrial 1,25-dihydroxyvitamin D3 24-hydroxylase mRNA, complete cds
AA007419	Human RGP4 mRNA, complete cds
T49657	Homo sapiens TWIK-related acid-sensitive K+ channel (TASK) mRNA, complete cds
N38959	Homo sapiens chaperonin containing t-complex polypeptide 1, beta subunit (Cctb) mRNA, complete cds
R51912	Human somatostatin I gene and flanks
H90415	Breast cancer 1, early onset
H41489	Adaptin, beta 1 (beta prime)
H15456	CALPAIN 1, LARGE
W45415	ELASTASE IIIB PRECURSOR
AA447751	Tyrosine hydroxylase
AA487486	Cyclin D1 (PRAD1; parathyroid adenomatosis 1)
R56604	Cholinergic receptor, nicotinic, alpha polypeptide 4
T65772	pulmonary surfactant protein (SP5)
H15085	ADP-ribosylation factor 4-like
R61295	Human ADP/ATP translocase mRNA, 3' end, clone pHAT8
T61256	H.sapiens KHK mRNA for ketohexokinase, clone pHHK3a
AA405731	Phosphoenolpyruvate carboxykinase 1 (soluble)
T71879	Complement component C2
R59927	Human mRNA for cytochrome c oxidase subunit VIc
AA496780	Human small GTP binding protein Rab7 mRNA, complete cds
AA176688	Human mRNA for lysosomal sialoglycoprotein, complete cds
AA436163	Homo sapiens Pig12 (PIG12) mRNA, complete cds
AA428778	Human placenta LERK-2 (EPLG2) mRNA, complete cds
AA463225	Bone morphogenetic protein 4
AA485426	Interferon (alpha, beta and omega) receptor 2
W47485	Human sigma receptor mRNA, complete cds
H84982	Human checkpoint suppressor 1 mRNA, complete cds
AA504615	Homo sapiens mRNA for CAB1, complete cds
H94487	Cathepsin E
AA448959	Homo sapiens NADH:ubiquinone oxidoreductase 15 kDa IP subunit mRNA, nuclear gene encoding mitochondrial protein, complete cds
AA070358	Transketolase (Wernicke-Korsakoff syndrome)
AA453401	Human PH-20 homolog (LUCA2) mRNA, partial cds
N66737	Collagen, type II, alpha 1 (primary osteoarthritis, spondyloepiphyseal dysplasia, congenital)
AA666180	Human v-erbA related ear-2 gene
AA857131	Human Tat-SF1 mRNA, complete cds
AA479102	Protein kinase C, beta 1

AA456077 Homo sapiens mRNA for p27, complete cds
 R87497 H.sapiens mRNA for 2.19 gene
 AA718910 Human tax1-binding protein TXBP181 mRNA, complete cds
 AA406269 Nuclear factor I/X (CCAAT-binding transcription factor)
 N74623 Insulin-like growth factor 2 (somatomedin A)
 H99364 Human chloride channel protein (CLCN7) mRNA, partial cds
 AA447684 Small proline-rich protein 1B (cornifin)
 AA282301 Homo sapiens nuclear dual-specificity phosphatase (SBF1) mRNA, partial cds
 H99588 Human lymphoid nuclear protein (LAF-4) mRNA, complete cds
 N53512 Homo sapiens alpha 2 delta calcium channel subunit isoform I mRNA, complete cds
 AA683321 Homo sapiens PAR-5 mRNA, probable 5' end
 AA608557 Damage-specific DNA binding protein 1 (127 kD)
 AA757764 Homo sapiens mRNA for DNA-binding protein, complete cds
 AA406064 Homo sapiens testis-specific Basic Protein Y 1 (BPY1) mRNA, complete cds
 N54596 Human Krueppel-related zinc finger protein (H-plk) mRNA, complete cds
 AA481988 Transcription factor 7 (T-cell specific)
 N62394 Gap junction protein, beta 1, 32kD (connexin 32, Charcot-Marie-Tooth neuropathy, X-linked)
 N26148 Zinc finger protein 148 (pHZ-52)
 AA496678 B-cell CLL/lymphoma 3
 AA400973 NEUTROPHIL GELATINASE-ASSOCIATED LIPOCALIN PRECURSOR
 AA497027 Human mRNA, clone HH109 (screened by the monoclonal antibody of insulin receptor substrate-1 (IRS-1))
 N64508 Homo sapiens podocalyxin-like protein mRNA, complete cds
 AA033564 H.sapiens mRNA for DGCR6 protein
 AA446108 Endoglin (Osler-Rendu-Weber syndrome 1)
 AA159577 Mucin 5, subtype B, tracheobronchial
 R36958 unknown EST
 AA629808 Ribosomal protein L6
 AA482067 Human tazarotene-induced gene 2 (TIG2) mRNA, complete cds
 AA669314 ATP synthase, H⁺ transporting, mitochondrial F1 complex, delta subunit
 AA775241 Aldolase A
 R73584 Homo sapiens hydroxysteroid sulfotransferase SULT2B1a (HSST2) mRNA, complete cds
 H28984 PHOSPHATIDYL SERINE SYNTHASE I
 R44202 Homo sapiens catechol-O-methyltransferase (COMT) mRNA, complete cds
 W70051 H.sapiens mRNA for M-phase phosphoprotein, mpp9
 AA401972 Human RalGDS-like 2 (RGL2) mRNA, partial cds
 AA236164 CATHEPSIN S PRECURSOR
 R22412 Platelet/endothelial cell adhesion molecule (CD31 antigen)
 AA424804 MULTIDRUG RESISTANCE-ASSOCIATED PROTEIN 1
 AA669443 Eukaryotic translation initiation factor 5 (eIF5)
 N69689 RAS-RELATED PROTEIN RAB-1A
 H24316 AQUAPORIN-CHIP
 AA074224 Recoverin
 R36571 Human U1 snRNP-specific protein A gene
 AA056465 Human 54 kDa protein mRNA, complete cds
 AA633811 H.sapiens E4BP4 gene
 AA457155 Human zinc-finger protein C2H2-150 mRNA, complete cds
 AA459104 60S RIBOSOMAL PROTEIN L13
 R40212 Human coatomer protein (HEPCOP) mRNA, complete cds
 AA086476 Adenosine monophosphate deaminase 1 (isoform M)
 AA663310 Thymidylate synthase
 AA455640 Homo sapiens signalosome subunit 3 (Sgn3) mRNA, complete cds
 AA496879 Human (clone E5.1) RNA-binding protein mRNA, complete cds

AA085749 Homo sapiens mRNA for ATP binding protein, complete cds
AA425755 Homo sapiens mRNA for leukemia associated gene 1
N52350 H.sapiens mRNA for protein-tyrosine-phosphatase (tissue type: testis)
AA630104 Lipase A, lysosomal acid, cholesterol esterase (Wolman disease)
AA454854 ALPHA-AMYLASE 2B PRECURSOR
W73406 DIHYDROPRYRIDINE-SENSITIVE L-TYPE, SKELETAL MUSCLE CALCIUM CHANNEL GAMMA SUBUNIT
R12802 Human cytochrome bc-1 complex core protein II mRNA, complete cds
AA465355 Homo sapiens mRNA for U3 snoRNP associated 55 kDa protein
AA829383 DUAL SPECIFICITY MITOGEN-ACTIVATED PROTEIN KINASE KINASE 3
AA629189 Keratin 4
AA430512 Homo sapiens cytoplasmic antiproteinase 3 (CAP3) mRNA, complete cds
AA456439 Human homozygous deletion target in pancreatic carcinoma (DPC4) mRNA, complete cds
H27864 SECRETOGRANIN II PRECURSOR
AA644657 MHC class I protein HLA-A (HLA-A28,-B40, -Cw3)
R40460 Homo sapiens phosphatidylinositol 4-kinase mRNA, complete cds
W96058 Human hnRNP H mRNA, complete cds
T72202 Human transcription factor IL-4 Stat mRNA, complete cds
AA598794 Connective tissue growth factor
AA599178 Ribosomal protein L27a
R88247 Adrenergic, beta, receptor kinase 1
T98612 Alpha-1 type 3 collagen
AA454856 Phospholipid hydroperoxide glutathione peroxidase
N67048 Type 3 iodothyronine deiodinase
AA778675 Homo sapiens mRNA for calmegin, complete cds
H51117 Human calmodulin dependent phosphodiesterase PDE1B1 mRNA, complete cds
N36174 5-HYDROXYTRYPTAMINE 2B RECEPTOR
AA777187 Homo sapiens Cyr61 mRNA, complete cds
R09561 Decay accelerating factor for complement (CD55, Cromer blood group system)
R16849 Human HsPex13p mRNA, complete cds
AA884167 ANNEXIN XIII
AA136983 Cadherin 11 (OB-cadherin)
AA488622 Human signal transducing adaptor molecule STAM mRNA, complete cds
AA699427 Fructose-bisphosphatase 1
AA490459 Transcobalamin II
AA626787 Human ras-related C3 botulinum toxin substrate (rac) mRNA, complete cds
N62179 Human methylmalonate semialdehyde dehydrogenase gene, complete cds
N27190 UBIQUITIN CARBOXYL-TERMINAL HYDROLASE ISOZYME L3
AA441895 Human glutathione-S-transferase homolog mRNA, complete cds
AA463924 FACTOR VIII INTRON 22 PROTEIN
N78843 Homo sapiens cyclophilin-33A (CYP-33) mRNA, complete cds
AA629719 Cytochrome c oxidase VIIc subunit
AA464755 Ankyrin 1, erythrocytic
AA459351 H.sapiens sds22-like mRNA
AA488346 MYOSIN LIGHT CHAIN ALKALI, SMOOTH-MUSCLE ISOFORM
AA427899 Human mRNA fragment encoding beta-tubulin. (from clone D-beta-1)
AA453813 H.sapiens mRNA for Gal-beta(1-3/1-4)GlcNAc alpha-2,3-sialyltransferase
AA397824 Dopachrome tautomerase (dopachrome delta-isomerase, tyrosine-related protein 2)
AA633901 Transforming growth factor, beta-induced, 68kD
AA181334 Troponin I (skeletal fast)
AA292410 Clusterin (complement lysis inhibitor; testosterone-repressed prostate message 2; apolipoprotein J)
AA253434 HEAT SHOCK FACTOR PROTEIN 2
AA455056 H.sapiens mRNA for MAP kinase activated protein kinase

R55188 Human pre-T/NK cell associated protein (3B3) mRNA, 3' end
 AA465723 Homo sapiens mRNA for protein phosphatase 2C gamma
 N49856 SODIUM- AND CHLORIDE-DEPENDENT BETAINE TRANSPORTER
 AA455272 H.sapiens mRNA for ITBA1 protein
 AA459292 CDC28 protein kinase 1
 AA878561 Ubiquitin A-52 residue ribosomal protein fusion product 1
 AA772066 Human phosphatidylinositol (4,5)bisphosphate 5-phosphatase homolog mRNA, partial cds
 N78621 H.sapiens mRNA for gamma-adaptin
 AA291490 H.sapiens mRNA for processing α -glucosidase I
 N46828 Homo sapiens mRNA for inositol 1,4,5-trisphosphate 3-kinase isoenzyme, partial cds
 AA150487 Alkaline phosphatase, placental (Regan isozyme)
 AA282537 MYOCYTE-SPECIFIC ENHANCER FACTOR 2
 AA707922 Human mRNA for cone-specific cGMP phosphodiesterase gamma subunit, complete cds
 AA443638 Homo sapiens breast cancer-specific protein 1 (BCSG1) mRNA, complete cds
 W73892 Human putative tumor suppressor (LUCA15) mRNA, complete cds
 N70734 Troponin T2 (cardiac)
 H57136 Human phospholemman chloride channel mRNA, complete cds
 AA709414 Nidogen (enactin)
 W65461 Human protein tyrosine phosphatase mRNA, complete cds
 AA436564 Human cellular proto-oncogene (c-mer) mRNA, complete cds
 AA029042 Human HSIAH2 mRNA, complete cds
 AA427725 Homo sapiens carboxypeptidase Z precursor, mRNA, complete cds
 N51280 ADP-ribosylation factor like 1
 AA281347 H.sapiens mRNA for MHC class I promoter binding protein
 AA402960 Human HLA class III region containing NOTCH4 gene, partial sequence, homeobox PBX2 (HPBX) gene, receptor for advanced glycosylation end products (RAGE) gene, complete cds, and 6 unidentified cds
 N98485 Human forkhead protein FREAC-2 mRNA, partial cds
 AA490209 H.sapiens mRNA for Sop2p-like protein
 W61361 Homo sapiens cytoplasmic antiproteinase 2 (CAP2) mRNA, complete cds
 N51018 Biglycan
 AA455281 DEFENDER AGAINST CELL DEATH 1
 W69471 V-ski avian sarcoma viral oncogene homolog
 AA486321 Vimentin
 AA458982 Solute carrier family 9 (sodium/hydrogen exchanger), isoform 1 (antiporter, Na+/H+, amiloride sensitive)
 AA442095 NEDD-4 PROTEIN
 N99003 Active BCR-related gene
 AA609284 Homo sapiens mRNA for Eph-family protein, complete cds
 AA195036 Human Ro/SSA ribonucleoprotein homolog (RoRet) mRNA, complete cds
 AA478268 Human CtBP mRNA, complete cds
 AA608583 Homo sapiens mRNA for OTK27, complete cds
 AA486435 Homo sapiens mRNA for CDEP, complete cds
 AA505045 Human L2-9 transcript of unarranged immunoglobulin V(H)5 pseudogene
 AA487893 TUMOR-ASSOCIATED ANTIGEN L6
 AA292226 Homo sapiens creatine transporter mRNA, complete cds
 H87106 Homo sapiens T245 protein (T245) mRNA, complete cds
 W96450 Human putative tRNA synthetase-like protein mRNA, complete cds
 N33331 Human peroxisome proliferator activated receptor mRNA, complete cds
 AA405800 Dodecenoyl-Coenzyme A delta isomerase (3,2 trans-enoyl-Coenzyme A isomerase)
 T51539 Macrophage stimulating 1 (hepatocyte growth factor-like)
 N59764 Human guanosine 5'-monophosphate synthase mRNA, complete cds
 AA521346 H.sapiens mRNA for Ndr protein kinase

AA428551 Homo sapiens SOX22 protein (SOX22) mRNA, complete cds
AA489383 Bone morphogenetic protein 2
AA490172 Collagen, type I, alpha-2
AA504477 Human cytoskeleton associated protein (CG22) mRNA, complete cds

List of Table III related sequences:

Accession_	NAME	Gene
mrgd		
M98539	Human prostaglandin D2 synthase gene, exon 7	
AB004922	Homo sapiens gene for Smad 3, exon 1, partial sequence	
AB006000	Homo sapiens mRNA for chondromodulin-1 precursor, complete cds	
AB017364	Homo sapiens mRNA for frizzled-2, complete cds	
AB020236	Homo sapiens gene for ribosomal protein L27A, complete cds	Ribosomal protein L27a
AB042820	Homo sapiens RPL6 gene for ribosomal protein L6, complete cds	Ribosomal protein L6
AB043547	Homo sapiens gene for SMAD4, partial cds	
AB080265	Homo sapiens CYP2J2 mRNA for cytochrome P450 2J2, complete cds	Cytochrome P450, subfamily 1I1 (arachidonic acid epoxygenase) polypeptide 2
AF000979	Homo sapiens testis-specific Basic Protein Y 1 (BPY1) mRNA, complete cds	Homo sapiens testis-specific Basic Protein Y 1 (BPY1) mRNA, complete cds
AF001450	Homo sapiens core binding factor alpha1 subunit (CBFA1) gene, exon 7 and complete cds	
AF004231	Homo sapiens monocyte/macrophage Ig-related receptor MIR-10 (MIR cl-10) mRNA, complete cds	Homo sapiens monocyte/macrophage Ig-related receptor MIR-10 (MIR cl-10) mRNA, complete cds
AF009801	Homo sapiens homeodomain protein (BAPX1) mRNA, complete cds	
AF010126	Homo sapiens breast cancer-specific protein 1 (BCSG1) mRNA, complete cds	Homo sapiens breast cancer-specific protein 1 (BCSG1) mRNA, complete cds
AF010316	Homo sapiens Pig12 (PIG12) mRNA, complete cds	Homo sapiens Pig12 (PIG12) mRNA, complete cds
AF013591	Homo sapiens homolog of the Aspergillus nidulans sudD gene product mRNA, complete cds	Homo sapiens homolog of the Aspergillus nidulans sudD gene product mRNA, complete cds
AF037204	Homo sapiens RING zinc finger protein (RZF) mRNA, complete cds	
AF043339	Homo sapiens macrophage inflammatory protein 1 alpha (MIP1a) mRNA, partial cds	
AF049656	Homo sapiens inducible nitric oxide synthase (iNOS) mRNA, complete cds	
AF072872	Homo sapiens frizzled 1 mRNA, complete cds	
AF188285	Homo sapiens bone morphogenetic protein 9 (BMP9) mRNA, complete cds	
AF189279	Homo sapiens group IIIE secretory phospholipase A2 mRNA, complete cds	
AF248634	Homo sapiens syndecan 3 (SDC3) mRNA, complete cds	
AF304431	Homo sapiens hypoxia-inducible factor 1 alpha subunit (HIF1A) mRNA, complete cds	

AF339054	Homo sapiens BCL2-associated X protein (BAX) gene, exons 1, 2 and partial cds	Ubiquitin A-52 residue ribosomal protein fusion product 1
AF348700	Homo sapiens ubiquitin A-52 residue ribosomal protein fusion product 1 (UBA52), mRNA, complete cds	
AF395008	Homo sapiens Interleukin 4 (IL4) gene, complete cds	
AF405705	Homo sapiens matrix metalloproteinase 3 (stromelysin 1, progelatinase) (MMP3) gene, complete cds	
AF411526	Homo sapiens nerve growth factor beta (NGFB) mRNA, complete cds	Nerve growth factor beta
AF469046	Homo sapiens macrophage migration inhibitory factor (MIF) mRNA, complete cds	
AF477981	Homo sapiens osterix mRNA, complete cds	
AJ279016	Homo sapiens mRNA for chondrocyte expressed protein 68 kDa (CEP-68 gene)	
AY043326	Homo sapiens keratin 4 (KRT4) gene, complete cds	Keratin 4
AY044847	Homo sapiens aggrecanase 1 (ADAMTS4) gene, complete cds	
D13748	Human mRNA for eukaryotic initiation factor 4A1	
D38255	Homo sapiens mRNA for CAB1, complete cds	
D45399	Human mRNA for cone-specific cGMP phosphodiesterase gamma subunit, complete cds	
D49738	Human cytoskeleton associated protein (CG22) mRNA, complete cds	
D49835	Homo sapiens mRNA for DNA-binding protein, complete cds	
D90040	Human mRNA for arylamine N-acetyltransferase (EC 2.3.1.5)	
J00306	Human somatostatin I gene and flanks	
J03191	Human profilin mRNA, complete cds	
J03592	Human ADP/ATP translocase mRNA, 3' end, clone pHAT8	
J04111	Human c-jun proto oncogene (JUN), complete cds, clone hcCJ-1	
J04177	Human alpha-1 type XI collagen (COL11A1) mRNA, complete cds	
J04973	Human cytochrome bc-1 complex core protein II mRNA, complete cds	
J05036	Human cathepsin E mRNA, complete cds	
K00065	Human superoxide dismutase (SOD-1) mRNA, complete cds	
K00650	Human fos proto-oncogene (c-fos), complete cds	
L05095	Homo sapiens ribosomal protein L30 mRNA, complete cds	
L08895	Homo sapiens MADS/MEF2-family transcription factor (MEF2C) mRNA, complete cds	60S RIBOSOMAL PROTEIN L30
L10347	Human pro-alpha1 type II collagen (COL2A1) gene exons 1-54, complete cds	
L11566	Homo sapiens ribosomal protein L18 (RPL18) mRNA, complete cds	60S RIBOSOMAL PROTEIN L18

L13286	Human mitochondrial 1,25-dihydroxyvitamin D3 24-hydroxylase mRNA, complete cds	Human mitochondrial 1,25-dihydroxyvitamin D3 24-hydroxylase mRNA, complete cds
L13463	Human helix-loop-helix basic phosphoprotein (GOS8) mRNA, complete cds	
L13616	Human focal adhesion kinase (FAK) mRNA, complete cds	
L13720	Homo sapiens growth-arrest-specific protein (gas) mRNA, complete cds	Homo sapiens growth-arrest-specific protein (gas) mRNA, complete cds
L22009	Human hnRNP H mRNA, complete cds	Human hnRNP H mRNA, complete cds
L28997	Homo sapiens ARL1 mRNA, complete cds	ADP-ribosylation factor like 1
L31409	Homo sapiens creatine transporter mRNA, complete cds	Homo sapiens creatine transporter mRNA, complete cds
L33930	Homo sapiens CD24 signal transducer mRNA, complete cds and 3' region	
L34059	Homo sapiens cadherin-4 mRNA, complete cds	
L41162	Homo sapiens collagen alpha 3 type IX (COL9A3) mRNA, complete cds	Creatine kinase B
L47647	Homo sapiens creatine kinase B mRNA, complete cds	Vimentin
M13994	Human B-cell leukemia/lymphoma 2 (bcl-2) proto-oncogene mRNA encoding bcl-2-alpha protein, complete cds	Human guanine nucleotide-binding protein G-s, alpha subunit mRNA, partial cds
M14144	Human vimentin gene, complete cds	Elastase 1, pancreatic (elastase IIa)
M14631	Human guanine nucleotide-binding protein G-s, alpha subunit mRNA, partial cds	
M16652	Human pancreatic elastase IIA mRNA, complete cds	
M20137	Human interleukin 3 (IL-3) mRNA, complete cds, clone pCD-SR-alpha	
M22636	Human U1 small nuclear ribonucleoprotein 70 kd protein mRNA, complete cds	U1 snRNP 70K protein
M37825	Human fibroblast growth factor-5 (FGF-5) mRNA, complete cds	
M57293	Human parathyroid hormone-related peptide (PTHRP) gene, exons 1A, 1B, 1C, and 2	Ribosomal protein S4, X-linked
M58458	Human ribosomal protein S4 (RPS4X) isoform mRNA, complete cds	Homo sapiens catechol-O-methyltransferase (COMT) mRNA, complete cds
M58525	Homo sapiens catechol-O-methyltransferase (COMT) mRNA, complete cds	
M58549	Human matrix Glα protein (MGP) mRNA, complete cds	Erythroid alpha-spectrin
M61877	Human erythroid alpha-spectrin (SPTA1) mRNA, complete cds	Insulin-like growth factor binding protein 6
M62402	Human insulin-like growth factor binding protein 6 (IGFBP-6) mRNA, complete cds	
M65062	Human insulin-like growth factor 2 (HSF2) mRNA, complete cds	HEAT SHOCK FACTOR PROTEIN 2
M65217	Human heat shock factor 2 (HSF2) mRNA, complete cds	
M76701	Homo sapiens zinc finger protein 35 (ZNF35) gene, exon 1	Tropomodulin
M77016	Human tropomodulin mRNA, complete cds	

M81768	Human Na/H antiporter (APNH1) mRNA, complete cds	Human Na/H antiporter (APNH1) mRNA, complete cds
M84489	Human extracellular signal-regulated kinase 2 mRNA, complete cds	Adenosine monophosphate deaminase (isoform E)
M84721	Human AMP deaminase (AMPD3) mRNA, complete cds	GALECTIN-2
M87842	Human S-lac lectin L-14-II (LGALS2) mRNA, complete cds	Connective tissue growth factor
M92934	Human connective tissue growth factor, complete cds	
M95610	Human alpha 2 type IX collagen (COL9A2) mRNA, partial cds	
M96684	H.sapiens Pur (pur-alpha) mRNA, complete cds	H.sapiens Pur (pur-alpha) mRNA, complete cds
M97676	Homo sapiens (region 7) homeobox protein (HOX7) mRNA, complete cds	
NM_0001194	Homo sapiens Hypoxanthine phosphoribosyltransferase 1 (Lesch-Nyhan syndrome) (HPRT1), mRNA	
NM_000213	Homo sapiens integrin, beta 4 (ITGB4), mRNA	
NM_000221	Homo sapiens ketohexokinase (fructokinase) (KHK), transcript variant a, mRNA	H.sapiens KHK mRNA for ketohexokinase, clone pHKH3a
NM_000235	Homo sapiens lipase A, lysosomal acid, cholesterol esterase (Wolman disease) (LIPA), mRNA	Lipase A, lysosomal acid, cholesterol esterase (Wolman disease)
NM_000358	Homo sapiens transforming growth factor, beta-induced, 68kD (TGFBI), mRNA	Transforming growth factor, beta-induced, 68kD
NM_000364	Homo sapiens troponin T2, cardiac (TNNT2), mRNA	Troponin T2 (cardiac)
NM_000537	Homo sapiens renin (REN), mRNA	RENIN PRECURSOR, RENAL
NM_000574	Homo sapiens decay accelerating factor for complement (CD55, Cromer group system) (DAF), mRNA	Decay accelerating factor for complement (CD55, Cromer blood group system)
NM_000600	Homo sapiens interleukin 6 (interferon, beta 2) (IL6), mRNA	
NM_000618	Homo sapiens insulin-like growth factor 1 (somatomedin C) (IGF1), mRNA	
NM_000632	Homo sapiens integrin, alpha M (complement component receptor 3, alpha; also known as CD11b (p170), macrophage antigen alpha polypeptide) (ITGAM), mRNA	Integrin, alpha M (complement component receptor 3, alpha; also known as CD11b (p170), macrophage antigen alpha polypeptide)
NM_000711	Homo sapiens bone gamma-carboxyglutamate (gla) protein (osteocalcin) (BGLAP), mRNA	BGLAP, mRNA
NM_000962	Homo sapiens prostaglandin-endoperoxide synthase 1 (prostaglandin G/H synthase and cyclooxygenase) (PTGS1), transcript variant 1, mRNA	Prostaglandin-endoperoxide synthase 1 (prostaglandin G/H synthase and cyclooxygenase)
NM_000977	Homo sapiens ribosomal protein L13 (RPL13), transcript variant 1, mRNA	60S RIBOSOMAL PROTEIN L13
NM_001012	Homo sapiens ribosomal protein L35a (RPL35A), mRNA	Ribosomal protein L35a
NM_001025	Homo sapiens ribosomal protein S8 (RPS8), mRNA	40S RIBOSOMAL PROTEIN S8
NM_001064	Homo sapiens ribosomal protein S23 (RPS23), mRNA	40S RIBOSOMAL PROTEIN S23
NM_001127	Homo sapiens transketolase (Wernicke-Korsakoff syndrome) (TKT), mRNA	Transketolase (Wernicke-Korsakoff syndrome)
		Adaptin, beta 1 (beta prime)

NM_001200	Homo sapiens bone morphogenetic protein 2 (BMP2), mRNA	AP1B1), mRNA
NM_001229	Homo sapiens caspase 9, apoptosis-related cysteine protease (CASP9), transcript variant alpha, mRNA	Bone morphogenic protein 2
NM_001511	Homo sapiens GRO1 oncogene (melanoma growth stimulating activity, alpha) (GRO1), mRNA	Transcript variant alpha, mRNA
NM_001565	Homo sapiens small inducible cytokine subfamily B (Cys-X-Cys), member 10 (SCYB10), mRNA	Interferon (gamma)-induced cell line; protein 10 from
NM_001632	Homo sapiens alkaline phosphatase, placental (Regan isozyme) (ALPP), mRNA	(Regan isozyme)
NM_001687	Homo sapiens ATP synthase, H ⁺ transporting, mitochondrial F1 complex, delta ATP synthase, H ⁺ transporting, mitochondrial F1 complex, delta subunit (ATP5D), mRNA	Alkaline phosphatase, placental (Regan isozyme)
NM_001718	Homo sapiens bone morphogenetic protein 6 (BMP6), mRNA	
NM_001745	Homo sapiens calcium modulating ligand (CAMLG), mRNA	Calcium modulating ligand
NM_001797	Homo sapiens cadherin 11, type 2, OB-cadherin (osteoblast) (CDH11), transcript variant 1, mRNA	
NM_001844	Homo sapiens collagen, type II, alpha 1 (primary osteoarthritis, spondyloepiphyseal dysplasia, congenital) (COL2A1), transcript variant 1, mRNA	(primary osteoarthritis, spondyloepiphyseal dysplasia, congenital)
NM_001912	Homo sapiens cathepsin L (CTSL), mRNA	
NM_001969	Homo sapiens eukaryotic translation initiation factor 5 (EIF5), mRNA	Eukaryotic translation initiation factor 5 (eIF5)
NM_002073	Homo sapiens guanine nucleotide binding protein (G protein), alpha z polypeptide (GNAZ), mRNA	Guanine nucleotide binding protein (G protein), alpha z polypeptide
NM_002094	Homo sapiens G1 to S phase transition 1 (GSPT1), mRNA	G1 to S phase transition 1
NM_002160	Homo sapiens hexabronchion (tenascin C, cytотactин) (HXB), mRNA	
NM_002211	Homo sapiens integrin, beta 1 (fibronectin receptor, beta polypeptide, antigen CD29 includes MDF2, MSK12) (ITGB1), mRNA	
NM_002379	Homo sapiens matrilin 1, cartilage matrix protein (MATN1), mRNA	
NM_002381	Homo sapiens matrilin 3 (MATN3) precursor, mRNA	
NM_002421	Homo sapiens matrix metalloproteinase 1 (interstitial collagenase) (MMP1), mRNA	
NM_002424	Homo sapiens matrix metalloproteinase 8 (neutrophil collagenase) (MMP8), mRNA	
NM_002427	Homo sapiens matrix metalloproteinase 13 (collagenase 3) (MMP13), mRNA	
NM_002591	Homo sapiens phosphoenolpyruvate carboxykinase 1 (soluble) (PCK1), mRNA	Phosphoenolpyruvate carboxykinase 1 (soluble)
NM_002619	Homo sapiens platelet factor 4 (PF4), mRNA	Platelet factor 4
NM_002722	Homo sapiens pancreatic polypeptide (PPY), mRNA	
NM_002738	Homo sapiens protein kinase C, beta 1 (PRKCB1), mRNA	Protein kinase C, beta 1
NM_002903	Homo sapiens recoverin (RCV1), mRNA	Recoverin

NM_003036	Homo sapiens v-ski sarcoma viral oncogene homolog (avian) (SKI), mRNA	V-ski avian sarcoma viral oncogene homolog
NM_003282	Homo sapiens troponin I, skeletal, fast (TNNI2), mRNA	Troponin I (skeletal fast)
NM_003385	Homo sapiens visinin-like 1 (VSNL1), mRNA	Visinin-like 1
NM_003395	Homo sapiens wingless-type MMTV integration site family, member 14 (WNT14), mRNA	
NM_004613	Homo sapiens transglutaminase 2 (C polypeptide, protein-glutamine-gamma-gamma-methyltransferase) (TGM2), mRNA	
NM_004832	Homo sapiens glutathione-S-transferase omega (GSTTLp28), mRNA	Human glutathione-S-transferase homolog mRNA, complete cds
NM_004994	Homo sapiens matrix metalloproteinase 9 (gelatinase B, 92kD gelatinase, 92kD type IV collagenase) (MMP9), mRNA	
NM_004995	Homo sapiens matrix metalloproteinase 14 (membrane-inserted) (MMP14), mRNA	
NM_005038	Homo sapiens peptidylprolyl isomerase D (cyclophilin D) (PPID), mRNA	40 KD PEPTIDYL-PROLYL CIS-TRANS ISOMERASE
NM_005186	Homo sapiens calpain 1, (m1/m2) large subunit (CAPN1), mRNA	CALPAIN 1, LARGE
NM_005346	Homo sapiens heat shock 70kD protein 1B (HSPA1B), mRNA	HEAT SHOCK 70 KD PROTEIN 1
NM_005438	Homo sapiens FOS-like antigen 1 (FOSL1), mRNA	
NM_005506	Homo sapiens CD36 antigen (collagen type I receptor, thrombospondin receptor)-like 2 (lysosomal integral membrane protein II) (CD36L2), mRNA	
NM_006289	Homo sapiens talin 1 (TLN1), mRNA	
NM_006988	Homo sapiens a disintegrin-like and metalloprotease (reprolysin type) with thrombospondin type 1 motif, 1 (ADAMTS1), mRNA	
NM_007306	Homo sapiens breast cancer 1, early onset (BRCA1), transcript variant	Breast cancer 1, early onset
NM_007352	Homo sapiens elastase 3B, pancreatic (ELA3B), mRNA	ELASTASE 3B PRECURSOR
NM_014000	Homo sapiens vinculin (VCL), transcript variant meta-VCL, mRNA	
NM_014470	Homo sapiens GTP-binding protein (RHO6), mRNA	
NM_018952	Homo sapiens homeo box B6 (HOXB6), mRNA	
NM_021019	Homo sapiens myosin, light polypeptide 6, alkali, smooth muscle and non-muscle (MYL6), transcript variant 1, mRNA	MYOSIN LIGHT CHAIN ALKALI, SMOOTH-MUSCLE ISOFORM
NM_033150	Homo sapiens collagen, type II, alpha 1 (primary osteoarthritis, spondyloepiphyseal dysplasia, congenital) (COL2A1), transcript variant 2, mRNA	
NM_053056	Homo sapiens cyclin D1 (PRAD1; parathyroid adenomatosis 1) (CCND1), mRNA	Cyclin D1 (PRAD1; parathyroid adenomatosis 1)
NM_080682	Homo sapiens vascular cell adhesion molecule 1 (VCAM1), transcript variant 2, mRNA	
S79854	Homo sapiens type 3 iodothyronine deiodinase mRNA, complete cds	Type 3 iodothyronine deiodinase
S83308	SOX5=Sry-related HMG box gene [alternatively spliced] [human, testis, mRNA, 1473 nt]	
U07424	Human putative tRNA synthetase-like protein mRNA, complete cds	

U07620	Human MAP kinase mRNA, complete cds	Human MAP kinase mRNA, complete cds
U08023	Human cellular proto-oncogene (c-mer) mRNA, complete cds	Human cellular proto-oncogene (c-mer) mRNA, complete cds
U09303	Human T cell leukemia L ERK-2 (EPLG2) mRNA, complete cds	
U09577	Homo sapiens lysosomal hyaluronidase (LUCA2/HYAL2) mRNA, complete cds	Human PH-20 homolog (LUCA2) mRNA, partial cds
U09825	Human acid finger protein mRNA, complete cds	Acid finger protein ZNF173
U13261	Homo sapiens eIF-2-associated p67 homolog mRNA, complete cds	Human eIF-2-associated p67 homolog mRNA, complete cds
U13660	Human cartilage-derived morphogenetic protein 1 (CDMP-1) mRNA, complete cds	
U13991	Human TATA-binding protein associated factor 30 kDa subunit (taffl30) mRNA, complete cds	Human TATA-binding protein associated factor 30 kDa subunit (taffl30) mRNA, complete cds
U14966	Human ribosomal protein L5 mRNA, complete cds	Ribosomal protein L5
U14971	Human ribosomal protein S9 mRNA, complete cds	
U15085	Human HLA-DMB mRNA, complete cds	Major histocompatibility complex, class II, DM beta
U16031	Human transcription factor IL-4 Stat mRNA, complete cds	Human transcription factor IL-4 Stat mRNA, complete cds
U16261	Human MDA-7 (mda-7) mRNA, complete cds	
U18299	Human damage-specific DNA binding protein DDBa p127 subunit (DDB1) mRNA, complete cds	Damage-specific DNA binding protein 1 (1127 kD)
U20980	Human chromatin assembly factor-I p60 subunit mRNA, complete cds	Human chromatin assembly factor-I p60 subunit mRNA, complete cds
U22409	Human parathyroid hormone/PTH-related peptide receptor (PTH/PTHRP) gene, exon 14 and complete cds	Human eukaryotic initiation factor 2B-epsilon mRNA, partial cds
U23028	Human eukaryotic initiation factor 2B-epsilon mRNA, partial cds	
U23946	Human putative tumor suppressor (LUCA15) mRNA, complete cds	Human putative tumor suppressor (LUCA15) mRNA, complete cds
U24152	Human p21-activated protein kinase (Pak1) gene, complete cds	Human protein kinase PAK1 mRNA, complete cds
U25789	Human ribosomal protein L21 mRNA, complete cds	Ribosomal protein L21
U27699	Human pephBG T-1 betaine-GABA transporter mRNA, complete cds	SODIUM- AND CHLORIDE-DEPENDENT BETAINE TRANSPORTER
U31202	Human noggin (NOGGIN) gene, complete cds, (NOG)	
U32169	Human pro-a2 chain of collagen type XI (COL11A2) gene, complete cds	Human p37NB mRNA, complete cds
U32907	Human p37NB mRNA, complete cds	Human tax1-binding protein TXBP181 mRNA, complete cds
U33822	Human tax1-binding protein TXBP181 mRNA, complete cds	Human cleavage and polyadenylation specificity factor mRNA, complete cds
U37012	Human cleavage and polyadenylation specificity factor mRNA, complete cds	Human zinc-finger protein C2H2-150 mRNA, complete cds
U38864	Human zinc-finger protein C2H2-150 mRNA, complete cds	

U40373	Human cell surface glycoprotein CD44 mRNA, complete cds	AQUAPORIN-CHIP
U41517	Human channel-like integral membrane protein (AQP-1) mRNA, clone AQP-1-1656, complete cds	
U43148	Human patched homolog (PTC) mRNA, complete cds	Friedreich ataxia
U43747	Human frataxin (FRDA) mRNA, complete cds	Bone morphogenetic protein 4
U43842	Homo sapiens bone morphogenetic protein-4 (hBMP-4) gene, complete cds	Human phosphatidylinositol (4,5)bisphosphate 5-
U45975	Human phosphatidylinositol (4,5)bisphosphate 5-phosphatase homolog mRNA, partial cds	phosphatase homolog mRNA, partial cds
U53204	Human plectin (PLEC1) mRNA, complete cds	Human plectin (PLEC1) mRNA, complete cds
U53347	Human neutral amino acid transporter B mRNA, complete cds	Human neutral amino acid transporter B mRNA, complete cds
U59289	Human H-cadherin mRNA, complete cds	
U59423	Human Smad1 mRNA, complete cds	Human osteoclast stimulating factor mRNA, complete cds
U63717	Homo sapiens osteoclast stimulating factor mRNA, complete cds	
U68723	Human checkpoint suppressor 1 mRNA, complete cds	
U70312	Homo sapiens integrin binding protein Del-1 (Del1) mRNA, complete cds	
U72245	Human phospholemman chloride channel mRNA, complete cds	Human phospholemman chloride channel mRNA, complete cds
U75283	Human sigma receptor mRNA, complete cds	Human sigma receptor mRNA, complete cds
U76992	Human Tat-SF1 mRNA, complete cds	Human Tat-SF1 mRNA, complete cds
U80998	Human basic helix-loop-helix DNA binding protein (TWIST) gene, complete cds	
U83460	Human high-affinity copper uptake protein (hCTR1) mRNA, complete cds	Human high-affinity copper uptake protein (hCTR1) mRNA, complete cds
U90547	Human Ro/SSA ribonucleoprotein homolog (RoRet) mRNA, complete cds	Human Ro/SSA ribonucleoprotein homolog (RoRet) mRNA, complete cds
U92268	Homo sapiens mitogen activated protein kinase p38-2 mRNA, complete cds	
U93181	Homo sapiens nuclear dual-specificity phosphatase (SBF1) mRNA, partial cds	Homo sapiens nuclear dual-specificity phosphatase (SBF1) mRNA, partial cds
X00129	Human mRNA for retinol binding protein (RBP)	
X00588	Human mRNA for precursor of epidermal growth factor receptor	
X02910	Human gene for tumor necrosis factor (TNF-alpha)	GELSOLIN PRECURSOR, PLASMA
X03742	Human gene for L apoferitin exons 1 and 2	Human mRNA for plasma gelsolin
X04412	Human mRNA for receptor of retinoic acid	Human mRNA for receptor of retinoic acid
X06614		

X12794	Human v-erbA related ear-2 gene	Human v-erbA related ear-2 gene
X14420	Human mRNA for pro-alpha-1 type 3 collagen	Human mRNA for pro-alpha-1 type 3 collagen
X51801	Human OP-1 mRNA for osteogenic protein	Human mRNA for alpha1(I)X collagen (long form)
X54412	Human mRNA for alpha1(I)X collagen (long form)	Homo sapiens mitochondrial coxII mRNA for cytochrome C oxidase II subunit
X55654	Homo sapiens mitochondrial coxII mRNA for cytochrome P-450 (11 Beta)	Cytochrome P450 11 beta
X55764	Human mRNA for cytochrome P-450 (11 Beta)	Human L2-9 transcript of unarranged immunoglobulin V(H)5 pseudogene
X58399	Human L2-9 transcript of unarranged immunoglobulin V(H)5 pseudogene	V(H)5 pseudogene
		Agammaglobulinaemia protein-tyrosine kinase atk
X58957	H.sapiens atk mRNA for agammaglobulinaemia tyrosine kinase	
X60188	Human ERK1 mRNA for protein serine/threonine kinase	
X60382	H.sapiens COL10A1 gene for collagen (alpha-1 type X)	
X67337	H.sapiens HPBRII-4 mRNA	
X70683	H.sapiens mRNA for SOX-4 protein	
X71661	H.sapiens ERGIC-53 mRNA	
X74795	H.sapiens P1-Cdc46 mRNA	
X76770	H.sapiens mRNA for glycerol kinase testis specific 2	
X78712	H.sapiens mRNA for processing a-glucosidase I	
X87237	H.sapiens mRNA for human giant larvae homolog	
X87342	H.sapiens mRNA for ITBA1 protein	
X92475	H.sapiens mRNA for VEGF-C protein	
X94216	Homo sapiens solute carrier family 16 (monocarboxylic acid transporters), member 1 (SLC16A1), mRNA	Solute carrier family 16 (monocarboxylic acid transporters), member 1
XM_001306	Homo sapiens adenosine monophosphate deaminase 1 (isoform M) (AMPD1), mRNA	Adenosine monophosphate deaminase 1 (isoform M)
XM_001316	Homo sapiens calponin 3, acidic (CNN3), mRNA	Calponin 3, acidic
XM_001324	Homo sapiens fibromodulin (FMOD), mRNA	
XM_001782	Homo sapiens alkaline phosphatase, liver/bone/kidney (ALPL), mRNA	Alkaline phosphatase, liver/bone/kidney
XM_001826	Homo sapiens glycican 1 (GPC1), mRNA	Glycan 1
XM_002321	Homo sapiens peroxisome proliferative activated receptor, gamma (PPARG), mRNA	
XM_003059	Homo sapiens catenin (cadherin-associated protein), beta 1 (88KD) (CTNNB1), mRNA	
XM_003222	Homo sapiens cytochrome c oxidase subunit VIIc (COX7C), mRNA	Cytochrome c oxidase VIIc subunit
XM_003730	Homo sapiens interleukin 3 (colony-stimulating factor, multiple) (IL3), mRNA	
XM_003752	Homo sapiens interleukin 3 (colony-stimulating factor, multiple) (IL3), mRNA	

XM_003913	Homo sapiens integrin, alpha 2 (CD49B, alpha 2 subunit of VLA-2 receptor) (ITGA2), mRNA
XM_004063	Homo sapiens early growth response 1 (EGFR1), mRNA
XM_006121	Homo sapiens cathepsin D (lysosomal aspartyl protease) (CTSD), mRNA
XM_009336	Homo sapiens cartilage oligomeric matrix protein (pseudoachondroplasia, epiphyseal dysplasia 1, multiple) (COMP), mRNA
XM_009915	Homo sapiens leukemia inhibitory factor (cholinergic differentiation factor) (LIF), mRNA
XM_010702	Homo sapiens cathepsin K (pyncodysostosis) (CTSK), mRNA
XM_012503	Homo sapiens matrix metalloproteinase 2 (gelatinase A, 72kD gelatinase, 72kD type IV collagenase) (MMP2), mRNA
XM_012651	Homo sapiens collagen, type I, alpha 1 (COL1A1), mRNA
XM_015434	Homo sapiens chitinase 3-like 1 (cartilage glycoprotein-39) (CHI3L1), mRNA
XM_016181	Homo sapiens wingless-type MMTV integration site family, member 5A (WNT5A), mRNA
XM_017096	Homo sapiens active BCR-related gene (ABR), mRNA
XM_017384	Homo sapiens matrix metalloproteinase 7 (matrilysin, uterine) (MMP7), mRNA
XM_017591	Homo sapiens annexin A6 (ANXA6), mRNA
XM_028204	Homo sapiens nuclear factor of kappa light polypeptide gene enhancer in B-cells 1 (p105) (NFKB1), mRNA
XM_028642	Homo sapiens integrin, alpha 5 (fibronectin receptor, alpha a polypeptide) (ITGA5), mRNA
XM_029245	Homo sapiens collagen, type I, alpha 2 (COL1A2), mRNA
XM_029796	Homo sapiens frizzled-related protein (FRZB), mRNA
XM_031221	Homo sapiens interleukin 1, alpha (IL1A), mRNA
XM_031288	Homo sapiens aggrecan 1 (chondroitin sulfate proteoglycan 1, large aggregating proteoglycan, antigen identified by monoclonal antibody A0122) (AGC1), mRNA
XM_031289	Homo sapiens interleukin 8 (IL8), mRNA
XM_032902	Homo sapiens integrin, alpha 1 (ITGA1), mRNA
XM_033470	Homo sapiens defender against cell death 1 (DAD1), mRNA
XM_033657	Homo sapiens heparan sulfate proteoglycan 2 (perlecan) (HSPG2), mRNA
XM_033878	Homo sapiens tissue inhibitor of metalloproteinase 1 (erythroid potentiating activity, collagenase inhibitor) (TIMP1), mRNA
XM_034023	Homo sapiens regulator of G-protein signalling 4 (RGS4), mRNA
XM_034556	Homo sapiens chloride channel 7 (CLCN7), mRNA
XM_034845	Homo sapiens phosphatase and tensin homolog (mutated in multiple advanced Phosphatase and tensin homolog (mutated in multiple cancers 1) (PTEN), mRNA
XM_034890	Homo sapiens fibrillin 1 (Marfan syndrome) (FBN1), mRNA
XM_035662	Homo sapiens cathepsin B (CTSB), mRNA
XM_035842	Homo sapiens small inducible cytokine A5 (RANTES) (SCYA5), mRNA
XM_036107	Homo sapiens integrin, beta 2 (antigen CD18 (p95), lymphocyte function-associated antigen 1; macrophage antigen 1 (mac-1) beta subunit)

(ITGB2), mRNA	Homo sapiens collagen, type XVIII, alpha 1 (COL18A1), mRNA	Collagen, type XVIII, alpha 1
XM_036175	Homo sapiens ATP binding protein associated with cell differentiation	Homo sapiens mRNA for ATP binding protein, complete cds
XM_037087	(APACD), mRNA	
XM_037646	Homo sapiens msx homeo box homolog 2 (Drosophila) (MSX2), mRNA	
XM_037965	Homo sapiens chondroadherin (CHAD), mRNA	
XM_038584	Homo sapiens tissue inhibitor of metalloproteinase 3 (Sorsby fundus dystrophy, pseudoinflammatory, pseudoinflammatory). (TIMP3), mRNA	
XM_039094	Homo sapiens SRY (sex determining region Y)-box 9 (campomelic dysplasia, autosomal sex-reversal) (SOX9), mRNA	
XM_040037	Homo sapiens adrenergic, beta, receptor kinase 1 (ADRBK1), mRNA	Adrenergic, beta, receptor kinase 1
XM_040385	Homo sapiens S-adenosylmethionine decarboxylase 1 (AMD1), mRNA	S-adenosylmethionine decarboxylase 1
XM_042153	Homo sapiens biglycan (BGN), mRNA	Biglycan
XM_042664	Homo sapiens nuclear autoantigenic sperm protein (histone-binding) (NASP), mRNA	Nuclear autoantigenic sperm protein (histone-binding)
XM_044120	Homo sapiens fibroblast growth factor receptor 3 (achondroplasia, thanatophoric dwarfism) (FGFR3), mRNA	
XM_045089	Homo sapiens ATPase, Cu++ transporting, beta polypeptide (Wilson disease) (ATP7B), mRNA	ATPase, Cu++ transporting, beta polypeptide (Wilson disease)
XM_045802	Homo sapiens paxillin (PXN), mRNA	
XM_045890	Homo sapiens ADP-ribosylation factor 4-like (ARF4L), mRNA	ADP-ribosylation factor 4-like
XM_045925	Homo sapiens decorin (DCN), mRNA	Decorin
XM_045926	Homo sapiens lumican (LUM), mRNA	
XM_046035	Homo sapiens integrin, alpha L (antigen CD11A (p180), lymphocyte function-associated antigen 1; alpha polypeptide) (ITGAL), mRNA	Integrin, alpha L (antigen CD11A (p180), lymphocyte function-associated antigen 1; alpha polypeptide)
XM_046758	Homo sapiens tensin (TNS), mRNA	
XM_046765	Homo sapiens thymidylate synthetase (TYMS), mRNA	Thymidylate synthase
XM_047231	Homo sapiens fibulin 1 (FBLN1), mRNA	
XM_047719	Homo sapiens transcription factor 7 (T-cell specific, HMG-box) (TCF7), mRNA	Transcription factor 7 (T-cell specific)
XM_047802	Homo sapiens a disintegrin-like and metalloprotease (reprolysin type) with thrombospondin type 1 motif, 5 (aggrecanase-2) (ADAMTS5), mRNA	with thrombospondin type 1 motif, 5 (aggrecanase-2) (ADAMTS5),
XM_048167	Homo sapiens troponin T1, skeletal, slow (TNNT1), mRNA	Troponin T1, skeletal, slow
XM_048201	Homo sapiens metallothionein 1L (MT1L), mRNA	Metallothionein 1L
XM_049177	Homo sapiens vascular endothelial growth factor B (VEGFB), mRNA	
XM_049518	Homo sapiens intercellular adhesion molecule 1 (CD54), human rhinovirus receptor (ICAM1), mRNA	Human rhinovirus Intercellular adhesion molecule 1 (CD54), human rhinovirus receptor

XM_049534	Homo sapiens amylase, alpha 2A; pancreatic (AMY2A), mRNA	Amylase, alpha 2A; pancreatic
XM_049690	Homo sapiens coatomer protein complex, subunit alpha (COPA), mRNA	Human coatomer protein (HEPCOP) mRNA, complete cds
XM_049864	Homo sapiens colony stimulating factor 3 (granulocyte) (CSF3), mRNA	
XM_049937	Homo sapiens insulin-like growth factor binding protein 4 (IGFBP4), mRNA	
XM_050846	Homo sapiens Indian hedgehog homolog (Drosophila) (IHH), mRNA	
XM_053809	Homo sapiens similar to chondroitin sulfate proteoglycan 2 (versican) (H. sapiens) (LOC153633), mRNA	
XM_054566	Homo sapiens collagen, type VI, alpha 1 (COL6A1), mRNA	
XM_054686	Homo sapiens caspase 3, apoptosis-related cysteine protease (CASP3), mRNA	
XM_055254	Homo sapiens fibronectin 1 (FN1), mRNA	
XM_058069	Homo sapiens matrix metalloproteinase 12 (macrophage elastase) (MMP12), mRNA	
XM_084239	Homo sapiens retinoic acid receptor responder (tazarotene induced) (RARRES2), mRNA	2 Human tazarotene-induced gene 2 (TIG2) mRNA, complete cds
XM_084263	Homo sapiens cytochrome c oxidase subunit V1c (COX6C), mRNA	Human mRNA for cytochrome c oxidase subunit V1c
XM_084285	Homo sapiens integral membrane protein 2A (ITM2A), mRNA	
XM_085705	Homo sapiens tissue inhibitor of metalloproteinase 2 (TIMP2), mRNA	
XM_086368	Homo sapiens MUF1 protein (MUF1), mRNA	
XM_096277	Homo sapiens collagen, type V, alpha 1 (COL5A1), mRNA	
Y00985	Human mRNA for manganese-containing superoxide dismutase	
Y07566	H.sapiens mRNA for RIT protein	H.sapiens mRNA for RIT protein
Y07570	H.sapiens mRNA for PHAPI2b protein	H.sapiens mRNA for PHAPI2b protein
Y08999	H.sapiens mRNA for Sop2p-like protein	H.sapiens mRNA for Sop2p-like protein
Y12692	Homo sapiens mRNA for WNT11 gene	
Y13936	Homo sapiens mRNA for protein phosphatase 2C gamma	
Y15227	Homo sapiens mRNA for leukemia associated gene 1	
Z22865	H.sapiens dermatopontin mRNA, complete CDS	Homo sapiens mRNA for leukemia associated gene 1
Z50781	H.sapiens mRNA for leucine zipper protein	
Z50853	H.sapiens mRNA for CLPP	

The current invention also encompasses the process of down compression of previously determined 467 genes to a lower number that is still able to characterize the desired number of different cellular status. At present, for the determination of 7 different cell types or 5 development stages, a minimum of 26 spots of different marker genes are preferred, much preferred about 200 such spots. For full information, at least one spot for each of the presently 467 genes (markers) is preferred. A reduction of spot number can be of relevance e.g. if under certain conditions only a small subset of those genes listed in Tab II is required for 10 analysis e.g. in clinical applications. This down compression can be achieved by determining the ratio of actual to target number of genes and then choosing from each cluster accordingly to the determined ratio the necessary number of genes to fulfill the requirement. This process requires to group the number of genes for each analysis of e.g. Tab. I into 15 representative cluster familys from where representative genes can be selected. Such clusters familys can be determined as shown in Figure 1, namely by grouping clusters together that show a similar expression pattern. For each cluster family a representative number of genes may be chosen according to the compression factor that has been defined. It can 20 easily be seen that for larger clusters like e.g. "A" in Tab I more genes are available to select while in other clusters like e.g. "E" in Tab I less are present. At the end of the process one needs to balance the procedure in order to preserve the characteristics of the expression profile. In order to do so the amount of genes for each analysis should at least be greater than 2 25 sequences or spots, respectively, of different genes and for the total array at least 30. In order to control such a process classical hierarchical clustering (Stanford) analysis can be performed and checked on graphical presentations like treeview (Stanford). Cluster analysis may group similar expression profiles in families and will allow distinguishing between different 30 cell sources and allows classification of these cell cultures (see Fig. 2). If the cell sources are not properly represented in the cluster analysis it means that the selected marker genes are not balanced.

Example of an cartilage specific micro array structure:

To produce a microarray with printed oligonucleotides sequences of approx. at least 10 mers, preferably at least 25 mers, some sequences of table II need to be further processed. Since some of the determined sequences in Tab II are only expressed sequence tags (herein referred to as EST), they do not correspond to or represent the full-length cDNA. Therefore the EST preferably is BLAST searched with the public database at NCBI and the corresponding full-length cDNA determined. Only by having the correct and full-length cDNA it is possible to design oligomeric sequences that are balanced to each other and minimize any cross reactivity. Exemplary polynucleotide sequences (targets) are provided in the sequence listing of Table III. The cartilage related polynucleotide sequences as e.g. listed in Table III and other polynucleotide sequences known as key cartilage genes from the literature can be immobilized on a substrate and used as hybridizable array elements in a microarray format. Such microarrays can be composed of a subset of oligonucleotides representing e.g. sequences listed on Tab. II but modified to represent only full-length cDNA sequences. The used polynucleotides for the production of such a microarray can either be 50mer or also PCR (polymerase chain reaction) products but at least need to be longer than 10 bases. It should be noted that for microarray production also PCR products from the corresponding determined sequences directly or the full length cDNA can be used and it is not restricted just to oligonucleotides.

Methods to anchor such oligonucleotides or polynucleotides on a solid support are described in literature, together with information on length dependent distances between each oligo or polynucleotides and spots . (see e.g. Principal and Practice, DNA microarrays: gene expression analysis B.Jordan, Springer, 2001)

When polynucleotides are employed as hybridizable array elements in a microarray and depending on the software used, the array elements may be organized in an ordered fashion so that each element is present at a specified location on the substrate. If the array elements are at specified locations on the substrate, the hybridization patterns and

intensities (which together create a unique expression profile) can be interpreted in terms of expression levels of particular genes. This expression profile can then be used and may be correlated with any effect associated with a tissue and/or compound or to be investigated with regard
5 to a specific tissue and/or compound and allows comparison with already existing data.

One of such useful application of using ordered polynucleotides on microarrays is e.g. the comparison of gene expression profiles from a new sample e.g. a tissue biopsy, with already determined
10 characteristic gene expression profiles that are preferably stored in a database. Such stored gene expression profiles are e.g. of major importance if microarrays are applied in the clinic. In this case advantageously a database is set up that stores the corresponding gene expression profiles and advantageously also all patient informations, e.g.
15 history, blood pressure etc. By including all patient data and gene expression profiles in the analysis process and then starting a comparison with an expression profile from a new biopsy, it becomes possible to achieve a stronger correlation with the clinical outcome. This will allow to determine which therapy shall be applied, or even to modify an existing
20 therapy, e.g. to add growth factor x at a concentration y during the *ex vivo* tissue engineering phase. It may also be the case that the biopsy sample will demonstrate a poor gene expression profile that precludes the successful application of a modern therapeutic cell/tissue approach. Such cases would then only qualify for traditional surgical approaches, and hence
25 would not obtain the benefits of the tissue engineering process.

In analogy, the assessment of *in vitro* produced cartilage can also be performed. In the same way as mentioned above cell culture parameters, like e.g. culture media conditions, growth factor concentration,
30 are preferably stored in a data base together with the corresponding gene expression profiles. Comparison of the database entry with new profiles of new samples can then be used to assess the quality of the new *in vitro* produced tissue.

Subject Arrays and their use:

It should be noted that the invention described here is not dependent on any special array format rather than the possibility to select from an extended list of 467 novel key cartilage genes as well as meaningful gene expression patterns. A presently preferred subject array is a novel cartilage specific microarray that includes 187 genes that in the scope of this invention have been determined to be cartilage related and 140 genes that have been connected to cartilage in literature (see also Tab III). Normally, in high-density array procedures up to 10000 genes are usually applied and are not specific for certain applications. As one major general drawback, this results in massive data overflow and impaired data analysis due to difficult data handling and procedures. A preferred array has in its current state a minimal number of 150 genes, presently much preferred at most 333 genes, all of those with demonstrated relevance within cartilage tissue. Another major limitation has become apparent. While the invention WO01/24833 A2 describes a few marker genes associated with cartilage phenotype stability they do not allow to extensively describe chondrocyte cultures in details. No comprehensive classification of the different cell populations and culture conditions is possible as well as no gene expression profile or fingerprint can be achieved. Gene expression profiles determined with a set of genes represented in Tab II may allow to perform a more comprehensive analysis of different cell cultures conditions. Furthermore it may allow to compare and classify different tissue or the result of the different applied cell culture conditions. The above mentioned topics may only be possible with the disclosed invention as outlined within the following applications.

The inventive array CART-CHIP 300™ may be applied to classify (quality control) any source material, such as human cartilage biopsies, mesenchymal stem cell containing bone marrow aspirate, or pre-chondrogenic cells containing tissue according to pre-defined categories with respect to their capacity to re-build or re-organize a hyaline cartilage-like matrix *in vitro*. A rough subdivision could be for example "A", "B", or "C". While "A" will easily produce cartilage-like matrix, "B" will require special treatment to achieve an implantable construct, and "C" will represent those

cases that do yet not qualify for such a procedure. This biopsy classification system will allow:

• Quality control of the starting biopsy material and therefore optimization of the downstream process regarding e.g. *in vitro* tissue engineering applications

• Diagnostic evaluation of the patient and candidate treatment methods (e.g. CARTIGRAFT™) to ensure a cost-optimized procedure

• Quality control of *in vitro* tissue engineered products

10 The subject array of the present invention can be employed for all kind of research and developmental studies related to *in vitro* tissue engineering of cartilage. The possibility to assess proliferation, differentiation or re-differentiation as well as de novo matrix formation processes through analyses and comparison of a plurality of key cartilage genes (positive/negative markers) within one single experiment replaces current trial and error approaches and is thus far more rational.

15 The subject array can be applied to screen all kind of drugs, e.g. hormones, growth factors, within *in vitro* chondrocyte cultures regarding a potential beneficial effect on proliferation, differentiation, de 20 novo matrix formation. The deduced expression profiles can then be compared with existing data of e.g. native cartilage tissue and used to further optimize the process. Additionally the expression profiles can be compared with data from human adult and human infant cartilage to deduce a pathway or a strategy of how to induce more tissue formation *in vitro*.

25 The subject array of the preferred embodiment is very well suitable to better understand reaction pathways leading to new responses of chondrocytes *in vitro*. Only key cartilage genes comprising the whole spectrum of functional gene categories are to be investigated. This can be used to study the complexity of degenerative cartilage process *in vitro* and 30 the respective influence of potential beneficial drugs.

The subject array may be used to optimize cultures for *in vitro* cartilage formation starting from human cell sources other than cartilage like e.g. mesenchymal stem cells or bone marrow aspirates.

This subject array will be preferably used as powerful alternative for conventional molecular biology tools beside more established histological and biochemical analyses. By focusing on the most prominent cartilage marker genes being either positive or negative, it is possible to 5 characterize cartilage or cartilage related tissues as well as cell cultures thereof. In this respect, the subject array can replace conventional RT-PCR studies performed to check for cartilage marker gene expression, e.g. collagen I versus collagen II, aggrecan versus versican. By applying this subject array the set of markers will be easily increased by simultaneously 10 simplifying the experimental procedure and enhancing the outcome.

The subject arrays of the present invention have several advantages compared to existing microarrays as well as to conventional gene expression tools such as RT-PCR, Northern Blots etc.

Most importantly, the subject arrays are all based on key 15 cartilage genes. Beyond all the key cartilage genes known from the literature (~100-200 genes), 467 additional cartilage relevant genes have been discovered. Thus a significantly increased pool of cartilage key genes exists to choose from for various applications. For instance, to understand degenerative processes as they occur in OA or RA by study of complex 20 biological reaction pathways, it is important to follow expression of a relatively large number of genes.

Examples

The examples are described for the purposes of illustration 25 and are not intended to limit the scope of the invention.

Example 1: Analysis of various human cartilage samples

Useful for characterizing chondrocyte cultures derived from different human cartilage samples (adult and fetal), where adult samples 30 are different with respect to their capacity to form living tissue engineered equivalents under high density culture conditions.

Adult chondrocytes show different gene expression clusters compared to fetal chondrocytes and can be further distinguished from samples that will not produce living cartilage constructs (failures).

Human chondrocytes from adult and fetal articular cartilage
5 were proliferated in DMEM-F12 medium containing 10% FCS over several passages and transferred to pellet cultures (0.5×10^6 cells) in serum free DMEM-F12 medium supplemented with Ascorbate and Insulin medium. Proliferated cells were directly lysed with RLT buffer (RNeasy® Mini Kit, Qiagen) after trypsin release from plastic substrate, shredded 10 (QIAshredder, Qiagen) and kept frozen at -80°C in lysis buffer for later processing. High density pellet cultures were cultivated for 2 weeks if not otherwise specified, subsequently washed with phosphate buffered saline (PBS) and lysed in RLT Buffer (supplied with RNeasy® Kit). Total RNA was isolated from all samples as described in the manual provided with the 15 RNeasy kit and stored at -80°C.

Fluorescent labeled aRNA (amplified RNA) constructs were obtained by *in vitro* reverse transcription of the RNA followed by an *in vitro* amplification reaction.

20 2 µg of isolated total RNA were used per sample to amplify RNA by applying only one cycle of *in vitro* transcription (IVT, Millenium Biologix AG, Application Note).

25 2 µg of total RNA from each sample was primed with oligo(dT)_{24-mer} (containing a T7 RNA Polymerase Promotor) and reverse transcribed using 400 Units SuperScript II reverse transcriptase enzyme, nucleotides, 5x Reaction Buffer and Dithiothreitol (DTT) as described in protocol provided with the enzyme. For ribonuclease protection 1 µL RNase inhibitor (10 Units) was used to prevent RNA degradation during first strand synthesis. This first 30 strand synthesis reaction was incubated for 1 hour at 42°C.

To the first strand synthesis reaction 93 µl nuclease free water, 30 µl second strand buffer (Invitrogen, Basel, Switzerland) and 1.5 µl nucleotide mix (dATP, dTTP, dGTP, dCTP, 25 mM each) was added.

Second strand synthesis reaction mix was obtained by
5 adding 40 Units E. coli polymerase I (New England Biolabs, BioConcept, Allschwil, Switzerland), 10 Units E. coli DNA Ligase (New England Biolabs, BioConcept, Allschwil, Switzerland) and 2.5 Units Ribonuclease H (Fermentas, Labforce AG, Nunningen, Switzerland). Reaction was incubated for 2 hours at 16°C.

10 After this incubation step remaining RNA was degraded by adding 7.5 µl 1M sodium hydroxid containing 2mM EDTA (Ethylenediaminetetraacetic acid) for 10 minutes at 65°C. 7.5 µl 1M Hydrochloric acid was added to neutralize the reaction.

The obtained double strand DNA was purified in a
15 QIAquick® PCR purification kit (Qiagen, Hilden, Germany) and concentrated to 7.5 µl. To this concentrated RNA following reagents were added to obtain aRNA synthesis mix: 2 µl ATP (Adenosine triphosphate, 75mM), 2 µl GTP (Cytidin triphosphate, 75mM), 2 µl GTP (Guanosin triphosphate, 75mM), 2 µl UTP (Uridin triphosphate, 75mM), 1.5 µl 5-(3-aminoallyl)-Uridin triphosphate and 2 µl reaction buffer and 2 µl Enzyme
20 mix (both provided with Ambion MegaScript Kit, Ambion, Cambridgeshire, United Kingdom).

This aRNA synthesis mix was incubated for 4 hours at 37°C. Remaining double strand DNA was digested by adding 1 µl Dnase I
25 for 15 min at 37°C. aRNA was cleaned and concentrated with an RNeasy® Mini Kit column (Qiagen, Hilden Germany) and then concentrated to a final volume of 9 µl.

Fluorescent dye molecules were coupled to the reactive aminoallyl groups of the incorporated a 5-(3-aminoallyl)-Uridin triphosphate
30 molecules. One aliquot of either Cy3™- or Cy5™-mono reactive dye (Amersham Biosciences, Buckinghamshire, United Kingdom) was diluted in 40 µl water free Dimethyl sulfonoxide. 10 µl of one of the diluted Cy™ mono reactive dyes was added to each sample buffered in 100mM Carbonate

buffer (pH 9.00). Reaction was quenched after 1 hour by adding 10.4 µl Ethanol amine for 15 min at room temperature.

Unincorporated dye molecules were removed by ethanol precipitation. 2 µl Glycogen (Invitrogen, Basel, Switzerland) was added as 5 carrier during precipitation. After precipitation aRNA pellet was washed with 80% ethanol, dried and resuspended in 50 µl 1x Fragmentation buffer (200mM Tris(hydroxymethyl)aminomethane hydrochloride, 500 mM Potassium acetate, 150mM Magnesium acetate). aRNA was fragmented for 35 min at 94°C and placed on ice immediately. Fragmented aRNA was 10 dissolved in 900 µl hybridization buffer.

For denaturation aRNA was incubated for 5 min at 98°C and centrifuged for 30 sec at full speed in a microcentrifuge.

One CART-CHIP™ 300 (Millenium Biologix AG, Switzerland) was placed face down in a standard hybridization chamber. 15 Hybridization solution containing the denatured and labeled aRNA sample was injected using a standard micropipet whereas Cy3™ and Cy5™ samples were hybridized together in one hybridization chamber (Millenium Biologix AG, Switzerland). The microarrays were incubated overnight at 42°C in a PCR thermal cycler (TGradient, Whatman Biometra GmbH, 20 Göttingen, Germany).

After incubation unspecific aRNA probe was washed away with 1xSSC, 0.1%SDS for 5 min at room temperature, followd by another wash step in 1x SSC, 0.1% SDS for 5 min and rinsed with 1xSSC without SDS for 1 minute to remove excessive SDS. 1xSSC was discarded. 25 Remaining 1xSSC buffer on the slide surface was removed by centrifuge the slide for 2 min at 1500 x g.

The dried CART-CHIP™ 300 were then scanned using an Affymetrix 418 microarray scanner.

Expression level raw data for every spot was obtained with ImageQuaNT 30 (Molecular Dynamics). Raw data was normalized by dividing every expression value by total expression value of all spots for every sample and filtered by setting all values below the 25 percentile to the value of this 25 percentile to remove noise (25 percentile threshold).

For each sample (e.g. de-differentiated and re-differentiated chondrocytes) a list of all measured genes was generated. This so called gene expression profile was then used for subsequent analyses.

Further data analysis was performed using either
5 hierarchical clustering with cluster.exe (written by Michael Eisen, Stanford University) or Self Organizing Maps (SOM), such as GeneCluster developed by Whitehead Institute (Massachusetts Institute of Technology, MIT). The settings of the software were optimized until a reasonable number of clusters resulted that were able to represent the comparison
10 thoroughly. In the following example the parameters were as following:

Basic parameters: SOM rows 6; SOM col:4; #epochs=3000;
#seeds=1

Advanced parameters: initialization: random vectors;
neighborhood: bubble; alpha l=2; alpha f=0.005; sigma l=3000; sigma f=2.

15 Fig 1 shows a typical result from a SOM analysis with the above mentioned basic parameters, whereas Fig 2 shows an example of a graphical presentation of a cluster analysis and viewed by the software treeview.

20 Example 2: Quality Control and Human Cartilage Sample

Classification

Useful to demonstrate how CART-CHIP™ 300 can be used to differentiate between diverse cell culture conditions, to distinguish different patients, to study the influence of 3D culture conditions and to
25 serve as a quality control tool during any tissue engineering process.

Human chondrocytes isolated from 4 different donors were proliferated over one passage (P1) and then cultivated as high density pellets (0.5×10^6 cells) in 3D culture for 7 and 14 days. RNA samples were taken from proliferated as well as from 3D cultured cells resulting in totally
30 12 different samples as shown in Figure 8. RNA isolated from this samples was shredded in a QIAshredder (QIAGEN, Hilden, Germany), amplified, hybridized, washed and scanned as described in Example 1.

Data sets for all 12 samples were extracted and normalized as described in Example 1 to perform cluster and SOM analysis as noted below. Cluster analysis was performed using normalized data computed with GeneCluster.

5 Fig 3 shows a picture of such a cluster analysis for all 12 samples (#1-#12) consisting of 20 clusters (c0-c19)

10 Every cluster represents a typical gene expression pattern for all 12 samples indicated by a point, starting from sample #1 on the left hand side to sample #12 on the right hand side in every cluster. For example cluster c0 represents the expression level of 104 genes in all 12 samples in a given range indicated by the lines located above and below the computed points.

15 Another example for gene expression levels that behave similar for different culture conditions and donors are depicted in clusters c3, c4, c9 and c10. Meaning that every subset of the three donor specific points #1-#3, #4-#6, #7-#9, #10-#12 (see Tab V for detailed description) have gene clusters that behave similar in all analyzed samples.

20 An example of differently behaving genes is indicated in cluster c13, representing 10 genes that behave similar in donor #1 and #2 but show a different gene expression patterns for donors #3 and #4.

More detailed analyses are shown in Fig 4, Fig 5, and Fig 6. The clusters produced in these figures clearly demonstrate differences as well as similarities in cell behavior for either t0, t7 or t14 days, respectively.

25 Another software algorithm that can be applied for analysis of large amounts of data coming from gene microarrays is called hierarchical cluster analysis, whereas genes and/or different conditions with similar behavior in gene expression are clustered together. All hierarchical cluster analyses were performed using Cluster software described in Eisen 30 et al. (1998) PNAS 95:14863) and displayed using treeview.exe developed by same author.

Fig 7 shows such a cluster of selected genes for all 12 samples analyzed. Every square is representing one single gene

expression value. Different intensity means different expression levels. Dark squares are representing samples without any significant change in gene expression compare to the other samples or patients. Bright squares are indicating samples in which genes are up- or down-regulated relative to 5 other samples analyzed. A so called cluster of genes is a group of genes that behave similar from one donor to the other donors.

Not only genes but also samples can be clustered together. These clusters are called similarity dendrograms, shown in the top part of Fig 7. These tree-like structures illustrates similarities in gene expression 10 between different samples or donors. The closer a sample (#1...#12) is located to another sample in this dendrogram the more similar gene pattern they have.

Interestingly to see is that the seven samples located at the right side of the dendrogram (samples #1, #2, #5, #7, #8, #10 and #11) are 15 clustered together. This samples are representing t0 and t7 conditions as described above (illustrated in Tab V), whereas a cluster of 4 samples in the middle of the dendrogram (samples #3, #6, #9 and #12) are representing only t14 samples. This means a microarray of the current invention is able to distinguish between de-differentiated, proliferated 20 samples (t0 and t7) and re-differentiated samples in a later stage (t14).

An outlier represents sample #4 located at the most left side of Fig 7. which represents proliferated chondrocytes (t0) from donor 2 and could not clustered together with the remaining proliferated samples. Interestingly, this sample that it is not similar to all other proliferated 25 samples (#1, #7 and #10) was impaired with its capacity to form cartilage tissue equivalents following expansion in 2D culture. The biochemical analysis revealed a lower amount of total collagen/DNA for this sample and immunohistochemistry with collagen II antibodies resulted in only weak staining for a collagen II.

Example 3: Aortic Fibroblasts vs. Chondrocytes

Example to differentiate between expanded chondrocytes and aortic fibroblasts cultivated over 14 days in 3D settings.

A human aortic fibroblast cell source was proliferated and
5 brought to 3D culture. RNA was isolated after 14 days of culture. Expression data analysis was performed as described in previous Examples 1 and 2 using CART-CHIP™ 300 microarray.

A hierarchical cluster analysis was performed as described in example 2. Samples representing 3D culture after 14 days (t14) were
10 included in said data analysis (samples #3, # 6, # 9 and #12, see Tab V).

The result of the described analysis can be seen in Fig 8. The upper part of the figure shows a dendrogram as described in example 2. Aortic fibroblasts are not clustered together with human chondrocytes. The cluster shows a significantly different pattern compared to all other
15 cultures.

Obviously a gene expression pattern of an aortic fibroblast cell source can be clearly separated from a gene expression pattern of human chondrocytes. A micorarray of the present invention is therefore not only able to study differences between different chondrocyte culture
20 conditions but also to distinguish between cells isolated from different tissues.

Example 4: Arthritic conditions vs. healthy conditions

Useful to distinguish between normal healthy chondrocyte
25 behavior from cells resembling an arthritic phenotype. Interleukin-1 β is known to play a central role in the inflammation and connective tissue destruction observed in both rheumatoid arthritis (RA) and osteoarthritis (OA). Stimulation of *in vitro* chondrocyte cultures with Interleukin-1 β thus represents a simple experimental arthritis model.

30 The chondrocyte cell source from donor 4 (see Tab V) was proliferated over 3 passages and then cultivated as high-density pellet cultures (0.5×10^6 cells) for 16 hours and 7 days either in the absence or presence of Interleukin-1 β (30 ng/mL). RNA was isolated from all samples,

hybridized to CART-CHIP™ 300 and expression profiles were generated as described in Example 1.

A hierarchical cluster analysis was performed as described in Example 1 and the dendrogram and a selection of the representative gene clusters are 5 shown in Fig 9. This clearly shows that already a short stimulus of Interleukin-1 β results in alteration of the chondrocyte phenotype with gene expression changes that can be distinguished from untreated normal chondrocyte cultures.

Appendix**Table I**

Experiment and correlated gene expression analysis	Number of marker genes for each experiment (analysis)	experiment (analysis)
2D marker adult vs. fetal/infant	151	A
2D /3D adult vs. fetal/infant	96	B
3D marker failure	165	C
3D marker adult/fetal/infant	350	D
2D/3D marker adult	48	E
Time dependent failure marker	75	F
3D failure marker	41	G
Apoptosis related failure markers	30	H

5

Table V

Sample Numbers (#)			
	Proliferation (t0)	7 days in 3D culture (t7)	14 days in 3D culture (t14)
Donor 1	#1	#2	#3
Donor 2	#4	#5	#6
Donor 3	#7	#8	#9
Donor 4	#10	#11	#12

10

Table IV shows the results of a bioinformatic analysis of gene expression profiles of the 467 cartilage specific marker genes.

H3 F4/17															
H3 F4/12d		H3 F4/10d		H3 F4/9d		H3 F4/8d		H3 F4/7d		H3 F4/6d		H3 F4/5d		H3 F4/4d	
Cluster		Activation		SOM Description		H1 V2 Edt		H1 V2 1d		H1 V2 3d		H1 V2 5d		H1 V2 7d	
AA25363	2	2D/3D	0.620149592	H5 P1 2D	H4 P1 2D	0.5436551567	0.4974926555	0.5295265207	0.971884748	3.56/02	7.00/32	0.322383185	0.4477391374	0.2577397886	0.4455137252
AA845168	2	2D/3D	1.822987384	H5 P2 2D	H4 P2 2D	0.5655012668	0.5436551567	2.134502741	2.105619189	1.2825492518	1.083104031	1.3842848489	1.047842418	0.2187593688	0.2132682422
R52549	22	2D/3D	42.15205285	H5 P3 2D	H4 P3 2D	0.5655012668	0.5436551565	38.210254471	24.222595655	23.28245333	22.051117875	25.57074708	22.051117875	25.57074708	26.82/025859
H77128	4	2D/3D	0.114952897	H5 P4 2D	H4 P4 2D	0.5655012668	0.5436551565	0.111618832	0.111618832	0.111618832	0.111618832	0.119151225	0.119151225	0.122945222	0.1194757937
AA845016	6	2D/3D	29.161521448	H5 P5 2D	H4 P5 2D	0.5655012668	0.5436551565	30.162024448	29.04204044	2.889872655	0.4526520231	0.384685951	0.458136505	1.498136505	0.4552568223
AA37895	6	2D/3D	20.841572293	H5 P6 2D	H4 P6 2D	0.5655012668	0.5436551565	20.841572293	11.265688485	11.734020554	11.655950851	11.6319378872	11.6319378872	11.6319378872	11.6319378872
AA846938	15	2D/3D	9.5571114182	H5 P7 2D	H4 P7 2D	0.5655012668	0.5436551565	13.653283438	14.057827259	10.865952024	10.865952024	10.865952024	10.865952024	10.865952024	10.865952024
AA846918	16	2D/3D	34.952552129	H5 P8 2D	H4 P8 2D	0.5655012668	0.5436551565	35.131930207	48.320573207	48.320573207	48.320573207	48.320573207	48.320573207	48.320573207	48.320573207
AA845457	16	2D/3D	6.235743244	H5 P9 2D	H4 P9 2D	0.5655012668	0.5436551565	58.320573207	59.103425261	44.422056759	44.422056759	44.422056759	44.422056759	44.422056759	44.422056759
AA872001	20	2D/3D	81.813983212	H5 P10 2D	H4 P10 2D	0.5655012668	0.5436551565	85.111884928	82.12028621	64.174988482	54.200154668	64.75076884	54.200154668	64.75076884	54.200154668
H95390	1	2D/3D	0.538758103	H5 P11 2D	H4 P11 2D	0.5655012668	0.5436551565	0.610970303	0.610970303	0.610970303	0.610970303	0.393057939	0.393057939	0.393057939	0.393057939
AA686276	11	2D/3D	0.538758103	H5 P12 2D	H4 P12 2D	0.5655012668	0.5436551565	1.230141115	1.140525329	1.140525329	1.140525329	1.140525329	1.140525329	1.140525329	1.140525329
AA408555	11	2D/3D	0.538758103	H5 P13 2D	H4 P13 2D	0.5655012668	0.5436551565	5.0119500555	5.072610576	4.193769533	4.193769533	4.193769533	4.193769533	4.193769533	4.193769533
H05220	2	2D/3D	0.287711454	H5 P14 2D	H4 P14 2D	0.5655012668	0.5436551565	0.573472051	0.573472051	0.573472051	0.573472051	0.216165158	0.216165158	0.216165158	0.216165158
W37766	3	2D/3D	1.237126762	H5 P15 2D	H4 P15 2D	0.5655012668	0.5436551565	1.244550879	1.244550879	1.244550879	1.244550879	1.016405889	1.016405889	1.016405889	1.016405889
AA873985	3	2D/3D	2.389357721	H5 P16 2D	H4 P16 2D	0.5655012668	0.5436551565	2.389357721	2.132327897	1.174416144	0.687688684	1.076137628	0.747123688	1.076137628	0.747123688
AA978890	3	2D/3D	2.975518143	H5 P17 2D	H4 P17 2D	0.5655012668	0.5436551565	2.975518143	2.3145254655	2.3145254655	2.3145254655	2.3145254655	2.3145254655	2.3145254655	2.3145254655
R54618	6	2D/3D	1.405764041	H5 P18 2D	H4 P18 2D	0.565320542	0.5435551402	1.028467036	1.028467036	1.028467036	1.028467036	0.5468545459	0.5468545459	0.5468545459	0.5468545459
AA458530	23	2D/3D	2D/3D failure vs. cartilage	H5 P19 2D	H4 P19 2D	0.5655012668	0.5436551565	1.237126762	1.237126762	1.237126762	1.237126762	1.237126762	1.237126762	1.237126762	1.237126762
W37684	11	2D/3D	2D/3D failure vs. cartilage	H5 P20 2D	H4 P20 2D	0.5655012668	0.5436551565	3.425240852	3.425240852	3.425240852	3.425240852	1.261056411	1.261056411	1.261056411	1.261056411
N63102	11	2D/3D	2D/3D failure vs. cartilage	H5 P21 2D	H4 P21 2D	0.5655012668	0.5436551565	1.863018187	1.863018187	1.863018187	1.863018187	1.058653717	1.058653717	1.058653717	1.058653717
R56871	14	2D/3D	3.059732027	H5 P22 2D	H4 P22 2D	0.5655012668	0.5436551565	2.459270294	2.31613535207	3.086160597	2.708681643	5.859716488	7.877144785	7.877144785	7.877144785
N81209	22	2D/3D	11.253165658	H5 P23 2D	H4 P23 2D	0.565304536	0.543504536	9.202602512	9.202602512	9.202602512	9.202602512	8.703162113	8.703162113	8.703162113	8.703162113
AA446559	16	2D/3D	2D/3D failure vs. cartilage	H5 P24 2D	H4 P24 2D	0.5655012668	0.5436551565	7.181120166	6.444659109	6.444659109	6.444659109	10.91912142	10.91912142	10.91912142	10.91912142
AA253588	11	2D/3D	2D/3D failure vs. cartilage	H5 P25 2D	H4 P25 2D	0.5655012668	0.5436551565	3.425240852	3.425240852	3.425240852	3.425240852	1.031040051	1.031040051	1.031040051	1.031040051
W37769	18	2D/3D	2D/3D failure vs. cartilage	H5 P26 2D	H4 P26 2D	0.5655012668	0.5436551565	2.562042945	2.562042945	2.562042945	2.562042945	1.016216598	1.016216598	1.016216598	1.016216598
AA21701	18	2D/3D	2D/3D failure (adult)	H5 P27 2D	H4 P27 2D	0.5655012668	0.5436551565	0.398515453	0.398515453	0.398515453	0.398515453	0.706770017	0.706770017	0.706770017	0.706770017
AA055987	17	2D/3D	2D/3D failure (adult)	H5 P28 2D	H4 P28 2D	0.5655012668	0.5436551565	7.971928216	7.971928216	7.971928216	7.971928216	0.007946934	0.007946934	0.007946934	0.007946934
AA84128	22	2D/3D	2D/3D failure (adult)	H5 P29 2D	H4 P29 2D	0.5655012668	0.5436551565	6.87008786	6.87008786	6.87008786	6.87008786	1.2301051617	1.2301051617	1.2301051617	1.2301051617
N26538	23	2D/3D	2D/3D failure (adult)	H5 P30 2D	H4 P30 2D	0.5655012668	0.5436551565	9.127538191	9.127538191	9.127538191	9.127538191	1.048650659	1.048650659	1.048650659	1.048650659
AA056933	23	2D/3D	2D/3D failure (adult)	H5 P31 2D	H4 P31 2D	0.5655012668	0.5436551565	7.502614849	7.502614849	7.502614849	7.502614849	0.705709017	0.705709017	0.705709017	0.705709017
AA055987	17	2D/3D	2D/3D failure (adult)	H5 P32 2D	H4 P32 2D	0.5655012668	0.5436551565	7.971928216	7.971928216	7.971928216	7.971928216	0.007946934	0.007946934	0.007946934	0.007946934
AA88182	4	2D/3D	2D/3D failure (adult)	H5 P33 2D	H4 P33 2D	0.5655012668	0.5436551565	1.159705334	1.159705334	1.159705334	1.159705334	1.030658307	1.030658307	1.030658307	1.030658307
H05730	23	2D/3D	2D/3D failure (adult)	H5 P34 2D	H4 P34 2D	0.5655012668	0.5436551565	1.161221576	1.161221576	1.161221576	1.161221576	0.687681649	0.687681649	0.687681649	0.687681649
AA655155	22	2D/3D	2D/3D failure (adult)	H5 P35 2D	H4 P35 2D	0.5655012668	0.5436551565	1.2013016167	1.2013016167	1.2013016167	1.2013016167	1.030658305	1.030658305	1.030658305	1.030658305
AA673551	1	2D/3D	2D/3D failure (adult)	H5 P36 2D	H4 P36 2D	0.5655012668	0.5436551565	0.715716774	0.715716774	0.715716774	0.715716774	0.687681648	0.687681648	0.687681648	0.687681648
H12230	1	2D/3D	2D/3D failure (adult)	H5 P37 2D	H4 P37 2D	0.5655012668	0.5436551565	0.687681648	0.687681648	0.687681648	0.687681648	0.687681648	0.687681648	0.687681648	0.687681648
AA655565	1	2D/3D	2D/3D failure (adult)	H5 P38 2D	H4 P38 2D	0.5655012668	0.5436551565	0.734937592	0.734937592	0.734937592	0.734937592	1.1506151648	1.1506151648	1.1506151648	1.1506151648
R45581	2	2D/3D	2D/3D failure (adult)	H5 P39 2D	H4 P39 2D	0.5655012668	0.5436551565	0.686230424	0.686230424	0.686230424	0.686230424	0.686230424	0.686230424	0.686230424	0.686230424
AA655398	22	2D/3D	2D/3D failure (adult)	H5 P40 2D	H4 P40 2D	0.5655012668	0.5436551565	0.686230424	0.686230424	0.686230424	0.686230424	0.686230424	0.686230424	0.686230424	0.686230424
N61025	4	2D/3D	2D/3D failure (adult)	H5 P41 2D	H4 P41 2D	0.5655012668	0.5436551565	0.686230424	0.686230424	0.686230424	0.686230424	0.686230424	0.686230424	0.686230424	0.686230424
T67270	5	2D/3D	2D/3D failure (adult)	H5 P42 2D	H4 P42 2D	0.5655012668	0.5436551565	1.2013016167	1.2013016167	1.2013016167	1.2013016167	0.686230424	0.686230424	0.686230424	0.686230424
AA77304	6	2D/3D	2D/3D failure (adult)	H5 P43 2D	H4 P43 2D	0.5655012668	0.5436551565	0.686230424	0.686230424	0.686230424	0.686230424	0.686230424	0.686230424	0.686230424	0.686230424
AA64743	7	2D/3D	2D/3D failure (adult)	H5 P44 2D	H4 P44 2D	0.5655012668	0.5436551565	1.4107632421	1.4107632421	1.4107632421	1.4107632421	0.686230424	0.686230424	0.686230424	0.686230424

AAB30408	8	2D/3D fabrik (adult)	1.633210695	1.578545169	1.256079761	1.036467957	13.3417746	1.024786523	0.853226594	1.049852404
AA6853050	6	2D/3D fabrik (adult)	1.604545745	1.683528526	1.276522622	1.013468535	15.81592721	1.56843476	0.487406114	0.058453728
AA775674	3	2D/3D fabrik (adult)	1.777534446	1.5601517	1.234779751	2.555684777	2.0465547476	5.143265562	0.571505161	0.355545251
AA22934	19	2D/3D fabrik (adult)	2.356263943	1.558468586	2.320262625	1.41155555	10.9261687	6.78282516	2.357213126	0.454463251
AA27397	21	3D adult vs. fetal	1.671961938	2.668127071	2.73507272	2.743130689	17.84127043	17.84127043	4.181657624	4.6167167177
AA28195	3	3D adult vs. fetal	2.071920814	2.869735256	2.723507272	7.48271081	5.35989208	5.632765104	2.2895676485	3.09255509
AA79724	3	3D adult vs. fetal	7.247587164	7.247587164	3.470387176	6.011264841	2.441015531	4.011719073	1.703591212	1.247492101
T40541	2	3D adult vs. fetal	12.31479553	3.28885946	1.494626573	2.621670024	1.84626573	6.610093772	0.545092957	1.063191634
N33214	2	3D adult vs. fetal	2.654733448	2.651878481	1.727115813	2.171595555	1.858545558	1.059412046	0.505412046	1.7186745094
WG399	2	3D adult vs. fetal	2.818657835	2.818657835	1.271595793	2.171595793	1.858545555	1.059412046	0.505412046	1.7186745094
HW545	2	3D adult vs. fetal	5.267765763	6.138577001	2.723507272	7.48271081	5.35989208	5.632765104	2.2895676485	3.09255509
T7124	2	3D adult vs. fetal	1.861657018	4.172040349	3.470387176	6.011264841	2.441015531	4.011719073	1.703591212	1.247492101
N95418	2	3D adult vs. fetal	3.21585913	1.588715156	1.572090076	1.624540483	1.593545145	1.188589385	0.458592957	0.545092957
AA20675	18	3D adult vs. fetal	20.406111687	13.61156773	23.352524913	19.0558124	22.705840494	16.6788312	29.35857379	32.48610018
AA682551	21	3D adult vs. fetal	17.51359555	2.818657835	2.818657835	1.271595793	2.171595793	1.858545555	1.059412046	0.505412046
AA27633	3	3D adult vs. fetal	2.589589566	1.888592091	1.244684015	1.6244684015	1.6244684015	1.059412046	0.505412046	1.7186745094
AA10288	21	3D adult vs. fetal	1.861657018	4.172040349	3.470387176	6.011264841	2.441015531	4.011719073	1.703591212	1.247492101
AA07697	20	3D adult vs. fetal	21.132250654	30.8385114	32.170712822	21.132250654	20.406111687	21.132250654	19.454575151	20.216748524
R27559	20	3D adult vs. fetal	13.515395255	13.515395255	26.829397423	21.450763633	18.959587825	16.959587825	35.3261308	32.48610018
N71528	20	3D adult vs. fetal	25.44239224	17.0952218	28.22129217	21.61152023	24.00871315	21.61152023	20.406111687	23.34221611
AA27633	3	3D adult vs. fetal	6.628562023	7.473747702	7.056577922	7.21722212	6.130314027	4.550770154	5.622778325	5.569254543
AA10288	20	3D adult vs. fetal	18.725154654	14.162285204	25.175181653	21.610155058	17.855052928	14.162285204	14.162285204	27.2055224
AA70273	19	3D adult vs. fetal	52.170521917	34.350704958	32.170712827	47.82452102	43.940405516	44.515410119	42.477813824	34.7657485
H05619	2	3D adult vs. fetal	18.959587825	18.959587825	18.959587825	18.959587825	18.959587825	18.959587825	18.959587825	20.216748524
AA05532	6	3D adult vs. fetal	18.323235906	18.052353906	18.659587825	17.70582218	17.70582218	17.70582218	17.70582218	17.70582218
AA10458	20	3D adult vs. fetal	6.628562023	7.473747702	7.056577922	7.21722212	6.130314027	4.550770154	5.622778325	5.569254543
AA04322	20	3D adult vs. fetal	12.613128224	10.835918224	11.753522755	12.613128224	10.835918224	11.753522755	10.835918224	20.2000016
AA70273	19	3D adult vs. fetal	52.170521917	34.350704958	32.170712827	47.82452102	43.940405516	44.515410119	42.477813824	34.7657485
NB619	2	3D adult vs. fetal	4.624203052	1.284836438	0.973905825	4.624203052	1.284836438	0.973905825	1.284836438	2.273577052
AA04052	6	3D adult vs. fetal	1.624203052	1.284836438	0.973905825	4.624203052	1.284836438	0.973905825	1.284836438	2.273577052
AA104749	5	3D adult vs. fetal	5.267765763	5.267765763	5.267765763	5.267765763	5.267765763	5.267765763	5.267765763	22.2979728
AA05334	5	3D adult vs. fetal	6.12613128224	6.12613128224	6.12613128224	6.12613128224	6.12613128224	6.12613128224	6.12613128224	11.95951447
R61501	6	3D adult vs. fetal	4.215197474	4.384462636	4.860976052	4.6292681636	4.6292681636	4.6292681636	4.6292681636	1.0357035228
NB651	6	3D adult vs. fetal	4.624203052	1.284836438	0.973905825	4.624203052	1.284836438	0.973905825	1.284836438	2.273577052
AA053748	5	3D adult vs. fetal	6.883534139	4.782974749	5.219581254	5.219581254	5.219581254	5.219581254	5.219581254	1.0357035228
AA01749	5	3D adult vs. fetal	5.267765763	5.267765763	5.267765763	5.267765763	5.267765763	5.267765763	5.267765763	22.2979728
AA045157	5	3D adult vs. fetal	3.705758259	4.12613128224	4.12613128224	4.12613128224	4.12613128224	4.12613128224	4.12613128224	11.95951447
AA43118	8	3D adult vs. fetal	4.314538152	4.215197474	4.384462636	4.860976052	4.6292681636	4.6292681636	4.6292681636	1.0357035228
N9239	3	3D adult vs. fetal	4.572213222	3.033519374	5.072023105	3.243795759	5.623795759	5.623795759	5.623795759	2.273577052
AA107718	5	3D adult vs. fetal	5.622458227	5.622458227	5.190162151	4.1313128224	4.1313128224	4.1313128224	4.1313128224	1.0357035228
AA2533413	3	3D adult vs. fetal	3.575321316	4.703758259	4.12613128224	4.12613128224	4.12613128224	4.12613128224	4.12613128224	2.273577052
AA046701	4	3D adult vs. fetal	7.2474740589	7.004203052	9.578771775	8.4247750272	9.578771775	10.342940503	11.033734755	11.033734755
AA184562	4	3D adult vs. fetal	7.627425507	6.217051711	6.217051711	6.217051711	6.217051711	6.217051711	6.217051711	11.95951447
AA468557	4	3D adult vs. fetal	6.417152683	6.975115261	10.422291719	7.967450727	8.417202154	8.417202154	8.417202154	1.0357035228
AA169742	4	3D adult vs. fetal	5.622121327	5.622458227	5.622458227	5.622458227	5.622458227	5.622458227	5.622458227	1.0357035228
AA059155	18	3D adult vs. fetal	21.324295632	19.069817533	30.01029002	23.958122024	20.8144049	33.012775159	28.226262594	3.061205108
AA539143	4	3D adult vs. fetal	7.615149711	5.735953945	5.134091911	6.221695938	4.722722656	5.375853938	6.142856213	3.4216851434
AA43226	4	3D adult vs. fetal	7.127244497	7.2474740589	9.04203052	8.425755232	10.072026592	11.753521351	12.227216703	10.501916714
AA458349	3	3D adult vs. fetal	6.462151512	6.462151512	6.462151512	6.462151512	6.462151512	6.462151512	6.462151512	2.15970081
AA650651	9	3D adult vs. fetal	5.625161512	5.625161512	5.625161512	5.625161512	5.625161512	5.625161512	5.625161512	7.871716277
AA107718	5	3D adult vs. fetal	5.622458227	5.622458227	5.622458227	5.622458227	5.622458227	5.622458227	5.622458227	17.037617178
AA2533413	3	3D adult vs. fetal	5.735953945	5.735953945	5.134091911	6.221695938	4.722722656	5.375853938	6.142856213	3.4216851434
AA046701	4	3D adult vs. fetal	7.127244497	7.2474740589	9.04203052	8.425755232	10.072026592	11.753521351	12.227216703	10.501916714
AA184562	4	3D adult vs. fetal	6.462151512	6.462151512	6.462151512	6.462151512	6.462151512	6.462151512	6.462151512	7.871716277
AA43118	8	3D adult vs. fetal	4.314538152	4.215197474	4.384462636	4.860976052	4.6292681636	4.6292681636	4.6292681636	1.0357035228
N9239	3	3D adult vs. fetal	5.622458227	5.622458227	5.622458227	5.622458227	5.622458227	5.622458227	5.622458227	1.0357035228
AA107718	5	3D adult vs. fetal	5.622458227	5.622458227	5.622458227	5.622458227	5.622458227	5.622458227	5.622458227	1.0357035228
AA2533413	3	3D adult vs. fetal	5.735953945	5.735953945	5.134091911	6.221695938	4.722722656	5.375853938	6.142856213	3.4216851434
AA046701	4	3D adult vs. fetal	7.127244497	7.2474740589	9.04203052	8.425755232	10.072026592	11.753521351	12.227216703	10.501916714
AA184562	4	3D adult vs. fetal	6.462151512	6.462151512	6.462151512	6.462151512	6.462151512	6.462151512	6.462151512	7.871716277
AA43118	8	3D adult vs. fetal	4.314538152	4.215197474	4.384462636	4.860976052	4.6292681636	4.6292681636	4.6292681636	1.0357035228
N9239	3	3D adult vs. fetal	5.622458227	5.622458227	5.622458227	5.622458227	5.622458227	5.622458227	5.622458227	1.0357035228
AA107718	5	3D adult vs. fetal	5.622458227	5.622458227	5.622458227	5.622458227	5.622458227	5.622458227	5.622458227	1.0357035228
AA2533413	3	3D adult vs. fetal	5.735953945	5.735953945	5.134091911	6.221695938	4.722722656	5.375853938	6.142856213	3.4216851434
AA046701	4	3D adult vs. fetal	7.127244497	7.2474740589	9.04203052	8.425755232	10.072026592	11.753521351	12.227216703	10.501916714
AA184562	4	3D adult vs. fetal	6.462151512	6.462151512	6.462151512	6.462151512	6.462151512	6.462151512	6.462151512	7.871716277
AA43118	8	3D adult vs. fetal	4.314538152	4.215197474	4.384462636	4.860976052	4.6292681636	4.6292681636	4.6292681636	1.0357035228
N923										

H65066 11	3D adult vs. failure	0.9322404085	19.08305972	1.3235240485	0.843205322	3.42676519	0.34276548	0.256322183	0.656615168	4.286907537	0.476165756	0.479705389	6.74E-02	1.020767749
AA158785 11	3D adult vs. failure	1.024544481	14.76220711	1.171923958	5.650846409	0.7854642166	1.050491022	1.020171532	1.020921018	1.020921018	1.020921018	1.020921018	1.020921018	1.942656569
AA465871 11	3D adult vs. failure	1.204024045	16.32124036	1.819684978	1.051476559	1.2435832378	4.85572561	0.587816889	11.845395163	2.157028817	0.265843105	0.265843105	0.265843105	1.303743017
T38411 14	3D adult vs. failure	1.21280482	14.88316906	1.77716274	0.65635165	0.5863231188	0.5863231188	0.5863231188	0.5863231188	0.5863231188	0.5863231188	0.5863231188	0.5863231188	1.580361835
R00835 20	3D adult vs. failure	4.116867612	2.768021652	2.545095883	2.768021652	0.5863231188	0.5863231188	0.5863231188	0.5863231188	0.5863231188	0.5863231188	0.5863231188	0.5863231188	0.5863231188
H65068 16	3D adult vs. failure	7.486261053	3.4202103988	3.917295183	3.205047795	4.021338911	7.438138134	2.107908251	2.786481372	1.276784815	1.529882688	1.784624277	2.741465653	1.963706377
H72228 14	3D adult vs. failure	1.122357249	10.877255229	1.107269259	0.894651168	0.693067551	5.481280912	0.181168944	0.025171213	0.0577472855	1.001641215	1.058341215	1.058341215	0.907756492
AA778177 14	3D adult vs. failure	0.813801868	11.38620018	0.884728519	0.739758542	0.624486657	3.553148409	0.348135461	0.500533332	4.336526268	4.438877031	0.328584239	0.342158325	1.848278914
N21578 14	3D adult vs. failure	0.485863698	0.382232944	0.485863698	0.485863698	0.485863698	0.485863698	0.485863698	0.485863698	0.485863698	0.485863698	0.485863698	0.485863698	0.485863698
AA00419 14	3D adult vs. failure	12.20218711	0.650768103	1.073720196	0.573720196	0.573720196	0.573720196	0.573720196	0.573720196	0.573720196	0.573720196	0.573720196	0.573720196	0.573720196
T40657 14	3D adult vs. failure	1.474835588	12.24545622	1.0585853458	1.158150176	4.72830076	3.039720458	4.037614043	4.037614043	1.234676181	0.125959577	3.27E-02	1.338393316	
N39859 14	3D adult vs. failure	0.656538518	12.21621754	1.222240119	0.390241865	3.677892922	7.55E-02	0.055622229	0.055622229	7.313895345	1.268404297	0.307595568	0.246991758	1.035716195
F61912 14	3D adult vs. failure	0.933165591	13.32686622	0.554065456	0.584709701	0.545970233	4.744685202	0.367556111	0.157082387	0.297490002	3.304783027	0.316505074	0.285956547	0.164570106
H93415 14	3D adult vs. failure	0.485863698	14.40639888	1.01501188	1.5435650847	0.485863698	0.485863698	0.485863698	0.485863698	0.485863698	0.485863698	0.485863698	0.485863698	0.485863698
AA07419 14	2D/3D adult vs. total	0.47385976	2.768021652	1.222351732	1.651727022	1.0585853458	1.0585853458	1.0585853458	1.0585853458	0.125959577	0.125959577	0.125959577	0.125959577	0.125959577
H14409 1	2D/3D adult vs. total	4.116867612	2.768021652	1.222351732	1.651727022	1.0585853458	1.0585853458	1.0585853458	1.0585853458	0.125959577	0.125959577	0.125959577	0.125959577	0.125959577
H18459 1	2D/3D adult vs. total	4.015465693	2.768021652	1.222351732	1.651727022	1.0585853458	1.0585853458	1.0585853458	1.0585853458	0.125959577	0.125959577	0.125959577	0.125959577	0.125959577
W45415 1	2D/3D adult vs. total	5.612236327	1.06521681	2.681703828	2.027058421	2.656897392	6.615895257	0.187478377	5.20E-02	0.722058962	0.692564765	0.1718687152	0.278888223	1.030511383
AA447761 1	2D/3D adult vs. total	3.707303455	4.789210773	2.222350567	2.437972776	2.631459156	7.62326101	0.763973734	0.524940441	0.524940441	0.524940441	0.524940441	0.524940441	0.524940441
AA487486 1	2D/3D adult vs. total	4.21356536	4.045765801	3.01623294	2.286202019	2.867529465	6.87529465	0.48478397	0.322923734	3.697172286	0.125959577	0.125959577	0.125959577	0.125959577
AA50604 1	2D/3D adult vs. total	1.416867612	2.768021652	1.222351732	1.651727022	1.0585853458	1.0585853458	1.0585853458	1.0585853458	0.125959577	0.125959577	0.125959577	0.125959577	0.125959577
T65772 1	2D/3D adult vs. total	4.015465693	2.768021652	1.222351732	1.651727022	1.0585853458	1.0585853458	1.0585853458	1.0585853458	0.125959577	0.125959577	0.125959577	0.125959577	0.125959577
H15205 1	2D/3D adult vs. total	4.632017688	2.552602568	1.8747482576	1.797824065	2.621610394	7.261610394	0.100168783	0.222509567	0.222509567	0.222509567	0.222509567	0.222509567	0.222509567
F81295 1	2D/3D adult vs. total	6.752262525	3.645420419	2.945152063	2.584538772	2.621610394	7.986612723	0.212944574	0.678529465	0.725E-02	0.686320155	0.161974764	0.394765632	1.055565615
T61256 1	2D/3D adult vs. total	5.2593434	3.045765801	3.01623294	1.626585706	2.423024955	6.87529465	0.7559177475	0.7559177475	0.7559177475	0.7559177475	0.7559177475	0.7559177475	0.7559177475
AA405731 0	2D/3D adult vs. total	3.467187567	2.974835588	2.974835588	2.974835588	2.974835588	2.974835588	2.974835588	2.974835588	2.974835588	2.974835588	2.974835588	2.974835588	2.974835588
T61789 1	2D/3D adult vs. total	4.193732869	4.21021681	2.535095883	2.535095883	2.535095883	2.535095883	2.535095883	2.535095883	2.535095883	2.535095883	2.535095883	2.535095883	2.535095883
R59227 1	2D/3D adult vs. total	4.822340777	4.93832171	1.797824065	1.8747482576	2.192202627	7.925477268	9.10E-02	0.073920395	8.920E-02	0.177597899	0.177597899	0.177597899	0.177597899
AA465789 0	2D/3D adult vs. total	1.514704484	28.444598443	1.135156078	0.912626215	0.912626215	0.912626215	0.912626215	0.912626215	0.912626215	0.912626215	0.912626215	0.912626215	0.912626215
AA485246 0	2D/3D adult vs. total	0.883613275	16.90708911	0.965095265	0.7559177475	2.393946025	1.062810388	0.7817878516	0.7817878516	0.7817878516	0.7817878516	0.7817878516	0.7817878516	0.7817878516
W47495 0	2D/3D adult vs. total	1.147451877	19.39019797	0.795067517	0.795067517	0.795067517	0.795067517	0.795067517	0.795067517	0.795067517	0.795067517	0.795067517	0.795067517	0.795067517
AA31613 0	2D/3D adult vs. total	1.416867612	4.21021681	2.535095883	2.535095883	2.535095883	2.535095883	2.535095883	2.535095883	2.535095883	2.535095883	2.535095883	2.535095883	2.535095883
AA29778 0	2D/3D adult vs. total	2.627059323	23.816162424	1.291565974	1.291565974	1.291565974	1.291565974	1.291565974	1.291565974	1.291565974	1.291565974	1.291565974	1.291565974	1.291565974
AA465225 0	2D/3D adult vs. total	1.845750443	22.470520543	1.542261155	1.416051919	1.282644596	1.076807945	0.207623068	0.207623068	0.207623068	0.207623068	0.207623068	0.207623068	0.207623068
AA485246 0	2D/3D adult vs. total	1.349593025	1.349593025	1.349593025	1.349593025	1.349593025	1.349593025	1.349593025	1.349593025	1.349593025	1.349593025	1.349593025	1.349593025	1.349593025
AA707038 2	2D/3D adult vs. total	0.812067475	1.745421971	1.504152379	1.384519446	1.275116851	1.0395450656	0.206452077	0.206452077	0.206452077	0.206452077	0.206452077	0.206452077	0.206452077
AA475010 1	2D/3D adult vs. total	2.731170292	3.625623025	1.2167739168	1.341583946	1.275116851	1.0395450656	0.206452077	0.206452077	0.206452077	0.206452077	0.206452077	0.206452077	0.206452077
R67497 7	2D/3D adult vs. total	2.705477038	4.518797607	2.115702505	2.145704043	1.870778446	1.488045485	1.0395450656	0.206452077	0.206452077	0.206452077	0.206452077	0.206452077	0.206452077
AA465810 9	2D/3D adult vs. total	1.7212414218	3.5458431737	1.824265244	1.3870421214	1.357781476	1.0395450656	0.206452077	0.206452077	0.206452077	0.206452077	0.206452077	0.206452077	0.206452077
AA465810 7	2D/3D adult vs. total	6.849777687	4.849370059	3.3765705139	2.027160704	1.636987474	1.0395450656	0.206452077	0.206452077	0.206452077	0.206452077	0.206452077	0.206452077	0.206452077
AA465810 9	2D/3D adult vs. total	1.454543975	29.42112597	0.638618073	0.50755939185	0.4232327456	0.4232327456	0.4232327456	0.4232327456	0.4232327456	0.4232327456	0.4232327456	0.4232327456	0.4232327456
N74623 4	2D/3D adult vs. total	0.739586231	1.0362053205	1.454465342	3.867045222	3.717165524	4.567652395	4.567652395	4.567652395	4.567652395	4.567652395	4.567652395	4.567652395	4.567652395
R69364 6	2D/3D adult vs. total	3.9774016745	3.9774016745	2.63202025	2.747673157	2.747673157	2.747673157	2.747673157	2.747673157	2.747673157	2.747673157	2.747673157	2.747673157	2.747673157
AA47684 6	2D/3D adult vs. total	0.42021681	0.42021681	0.42021681	0.42021681	0.42021681	0.42021681	0.42021681	0.42021681	0.42021681	0.42021681	0.42021681	0.42021681	0.42021681
AQ482201 0	2D/3D adult vs. total	1.747161704	1.747161704	1.747161704	1.747161704	1.747161704	1.747161704	1.747161704	1.747161704	1.747161704	1.747161704	1.747161704	1.747161704	1.747161704
H95585 5	2D/3D adult vs. total	1.												

AAG65821	6	2D/3D adult vs. fetal	1.132326265	0.9683128368	0.71972912	0.246569776	0.293262449	0.563371987	0.123251357	0.2226202587	0.1812828534	1.446843052	
AAG65857	5	2D/3D adult vs. fetal	2.577273943	2.807387159	2.022820214	1.932689782	0.478935818	0.3859738555	0.926202116	0.310230768	0.238678638	1.726847773	
AAJ7774	2	2D/3D adult vs. fetal	6.650914196	3.357895765	3.861101866	2.752078921	2.955536228	0.043391314	0.297373976	0.165726525	0.1454573768	0.986769324	
AA46564	2	2D/3D adult vs. fetal	6.957105219	3.655696449	3.922727479	3.833308615	5.142577569	0.246550235	0.963197456	0.184985842	0.778897405	0.3025523413	
N4596	20	2D/3D adult vs. fetal	6.411654414	12.470210209	12.602761075	11.876528458	11.876528458	0.020015252	0.187903438	0.2217807313	0.16296158	0.210150557	23.7826774
AA40180	7	2D/3D adult vs. fetal	6.762916769	7.616281816	7.360134597	3.598121217	12.026444853	0.523630389	0.2922745158	0.1445465158	0.162961567	0.570859423	1.454573757
N2294	3	2D Marker adult vs. fetal	1.407716704	4.457515241	4.119161688	3.689785978	3.689785978	0.045716045	0.232745158	0.1919547638	0.1813619638	0.190205885	0.2075979351
N2048	3	2D Marker adult vs. fetal	3.44985341	3.44985341	3.44985341	3.44985341	3.142376907	0.040464813	0.626264957	0.2323240313	0.118160173	0.634595551	0.465656326
AA46570	3	2D Marker adult vs. fetal	7.00110843	4.055685653	5.257405653	5.722236323	5.160847823	0.101822059	0.319441345	0.257767754	0.178071395	0.156592754	1.173221765
AA40053	3	2D Marker adult vs. fetal	2.893464833	1.470210209	1.842847841	1.842847841	1.842847841	0.020015252	0.187903438	0.2217807313	0.16296158	0.210150557	20.422646267
AA40057	3	2D Marker adult vs. fetal	1.302515483	0.576716212	1.289729562	1.289729562	1.289729562	0.020015252	0.187903438	0.2217807313	0.16296158	0.210150557	1.61094171
AA40727	3	2D Marker adult vs. fetal	1.382281583	1.382281583	1.382281583	1.382281583	1.521052047	0.020015252	0.187903438	0.2217807313	0.16296158	0.210150557	2.675970931
NC4596	3	2D Marker adult vs. fetal	3.276856184	3.276856184	3.276856184	3.276856184	3.276856184	0.020015252	0.187903438	0.2217807313	0.16296158	0.210150557	1.173221765
AA02354	3	2D Marker adult vs. fetal	3.131882612	3.131882612	3.131882612	3.131882612	3.131882612	0.020015252	0.187903438	0.2217807313	0.16296158	0.210150557	1.258329556
AA44610	3	2D Marker adult vs. fetal	2.078301835	2.211010494	2.708301835	2.708301835	2.708301835	0.020015252	0.187903438	0.2217807313	0.16296158	0.210150557	0.840188206
AA40057	3	2D Marker adult vs. fetal	2.893464833	1.470210209	1.842847841	1.842847841	1.842847841	0.020015252	0.187903438	0.2217807313	0.16296158	0.210150557	1.95574052
AA40057	3	2D Marker adult vs. fetal	3.400658977	3.400658977	3.400658977	3.400658977	3.400658977	0.020015252	0.187903438	0.2217807313	0.16296158	0.210150557	1.454573757
AA40058	3	2D Marker adult vs. fetal	1.749559144	1.749559144	1.749559144	1.749559144	1.749559144	0.020015252	0.187903438	0.2217807313	0.16296158	0.210150557	1.042280575
AA62010	5	2D Marker adult vs. fetal	1.920165597	0.7905096	1.215971675	1.215971675	1.215971675	0.020015252	0.187903438	0.2217807313	0.16296158	0.210150557	1.148267897
AA44207	3	2D Marker adult vs. fetal	1.502353047	3.823255324	3.823255324	3.823255324	3.823255324	0.020015252	0.187903438	0.2217807313	0.16296158	0.210150557	0.595055889
AA465914	3	2D Marker adult vs. fetal	6.809536569	4.133265612	5.322740251	5.322740251	5.322740251	0.020015252	0.187903438	0.2217807313	0.16296158	0.210150557	0.1937273233
AA77541	3	2D Marker adult vs. fetal	4.651654793	5.624749503	5.624749503	5.624749503	5.624749503	0.020015252	0.187903438	0.2217807313	0.16296158	0.210150557	0.1937273233
AA77541	3	2D Marker adult vs. fetal	4.749559144	5.624749503	5.624749503	5.624749503	5.624749503	0.020015252	0.187903438	0.2217807313	0.16296158	0.210150557	0.1937273233
H2984	5	2D Marker adult vs. fetal	1.741764648	1.741764648	1.741764648	1.741764648	1.741764648	0.020015252	0.187903438	0.2217807313	0.16296158	0.210150557	0.1937273233
R4402	3	2D Marker adult vs. fetal	3.131882612	3.131882612	3.131882612	3.131882612	3.131882612	0.020015252	0.187903438	0.2217807313	0.16296158	0.210150557	0.1937273233
W7051	4	2D Marker adult vs. fetal	4.649745555	4.649745555	4.649745555	4.649745555	4.649745555	0.020015252	0.187903438	0.2217807313	0.16296158	0.210150557	0.1937273233
AA40192	5	2D Marker adult vs. fetal	4.222620244	5.237916385	5.237916385	5.237916385	5.237916385	0.020015252	0.187903438	0.2217807313	0.16296158	0.210150557	0.1937273233
AA21614	5	2D Marker adult vs. fetal	5.870131020	5.870131020	5.870131020	5.870131020	5.870131020	0.020015252	0.187903438	0.2217807313	0.16296158	0.210150557	0.1937273233
R22112	5	2D Marker adult vs. fetal	3.222360737	3.822360737	3.822360737	3.822360737	3.822360737	0.020015252	0.187903438	0.2217807313	0.16296158	0.210150557	0.1937273233
AA42404	4	2D Marker adult vs. fetal	6.723361642	6.844704705	6.844704705	6.844704705	7.010333442	7.630735938	7.630735938	7.630735938	7.630735938	10.780004633	1.0304297022
AA65949	5	2D Marker adult vs. fetal	3.473141273	6.646240449	6.646240449	6.646240449	6.646240449	4.619190227	4.619190227	4.619190227	4.619190227	19.07682409	30.085816229
N6989	2	2D Marker adult vs. fetal	4.649526593	5.237916385	5.237916385	5.237916385	5.237916385	0.020015252	0.187903438	0.2217807313	0.16296158	0.210150557	0.1937273233
AA21614	4	2D Marker adult vs. fetal	4.174728977	5.671745830	5.671745830	5.671745830	5.671745830	0.020015252	0.187903438	0.2217807313	0.16296158	0.210150557	0.1937273233
A07424	3	2D Marker adult vs. fetal	4.337835704	4.287835704	4.287835704	4.287835704	4.287835704	0.020015252	0.187903438	0.2217807313	0.16296158	0.210150557	0.1937273233
F3571	4	2D Marker adult vs. fetal	6.723361642	6.844704705	6.844704705	6.844704705	6.844704705	0.020015252	0.187903438	0.2217807313	0.16296158	0.210150557	0.1937273233
AA05845	4	2D Marker adult vs. fetal	5.626585245	6.057970025	6.057970025	6.057970025	6.057970025	0.020015252	0.187903438	0.2217807313	0.16296158	0.210150557	0.1937273233
AAG65811	4	2D Marker adult vs. fetal	5.047585747	5.641758567	5.641758567	5.641758567	5.641758567	0.020015252	0.187903438	0.2217807313	0.16296158	0.210150557	0.1937273233
AA45165	4	2D Marker adult vs. fetal	4.056853587	4.570928987	4.570928987	4.570928987	4.570928987	0.020015252	0.187903438	0.2217807313	0.16296158	0.210150557	0.1937273233
AA45165	4	2D Marker adult vs. fetal	2.916840447	4.249717776	4.249717776	4.249717776	4.249717776	0.020015252	0.187903438	0.2217807313	0.16296158	0.210150557	0.1937273233
R44122	4	2D Marker adult vs. fetal	6.565605623	6.565605623	6.565605623	6.565605623	6.565605623	0.020015252	0.187903438	0.2217807313	0.16296158	0.210150557	0.1937273233
AAB85476	4	2D Marker adult vs. fetal	3.026168534	3.238236469	3.238236469	3.238236469	3.238236469	0.020015252	0.187903438	0.2217807313	0.16296158	0.210150557	0.1937273233
AA658210	5	2D Marker adult vs. fetal	2.169342225	2.651352455	2.651352455	2.651352455	2.651352455	0.020015252	0.187903438	0.2217807313	0.16296158	0.210150557	0.1937273233
AAB2104	1	2D Marker adult vs. fetal	1.689359637	1.689359637	1.689359637	1.689359637	1.689359637	0.020015252	0.187903438	0.2217807313	0.16296158	0.210150557	0.1937273233
AAB2104	1	2D Marker adult vs. fetal	0.627874118	0.621602027	0.621602027	0.621602027	0.621602027	0.020015252	0.187903438	0.2217807313	0.16296158	0.210150557	0.1937273233
AAB2104	1	2D Marker adult vs. fetal	0.627874118	0.621602027	0.621602027	0.621602027	0.621602027	0.020015252	0.187903438	0.2217807313	0.16296158	0.210150557	0.1937273233
W73406	2	2D Marker adult vs. fetal	1.134525157	1.145219429	1.145219429	1.145219429	1.145219429	0.020015252	0.187903438	0.2217807313	0.16296158	0.210150557	0.1937273233
R12502	2	2D Marker adult vs. fetal	2.169342225	2.651352455	2.651352455	2.651352455	2.651352455	0.020015252	0.187903438	0.2217807313	0.16296158	0.210150557	0.1937273233
N5259	1	2D Marker adult vs. fetal	1.247318987	1.247318987	1.247318987	1.247318987	1.247318987	0.020015252	0.187903438	0.2217807313	0.16296158	0.210150557	0.1937273233
AAB32104	1	2D Marker adult vs. fetal	1.689359637	1.689359637	1.689359637	1.689359637	1.689359637	0.020015252	0.187903438	0.2217807313	0.16296158	0.210150557	0.1937273233
AAB32104	1	2D Marker adult vs. fetal	0.627874118	0.621602027	0.621602027	0.621602027	0.621602027	0.020015252	0.187903438	0.2217807313	0.16296158	0.210150557	0.1937273233
AAB32104	1	2D Marker adult vs. fetal	0.627874118	0.621602027	0.621602027	0.621602027	0.621602027	0.020015252	0.187903438	0.2217807313	0.16296158	0.210150557	0.1937273233
AAB32104	1	2D Marker adult vs. fetal	0.627874118	0.621602027	0.621602027	0.621602027	0.621602027	0.020015252	0.187903438	0.2217807313	0.16296158	0.210150557	0.1937273233
AAB32104	1	2D Marker adult vs											

H2764	0	2D Marker adult vs. fetal	4.0830675739	3.247167731	3.6823939494	9.780274543	10.79728458	7.985471283	7.17120518
AA84457	0	2D Marker adult vs. fetal	2.7495567533	1.946526953	2.8353883935	2.282715575	2.77240442	14.472886784	15.19475588
R40460	0	2D Marker adult vs. fetal	2.617744205	2.770494985	2.530348837	2.347568897	2.6319132034	13.67189221	8.252597769
W6505	1	2D Marker adult vs. fetal	0.71825179353	0.71825179355	0.715948884	0.647054285	0.5114066355	9.242656521	6.561662584
T7226	2	2D Marker adult vs. fetal	1.3721233229	0.954777756	1.233822319	1.209148557	1.122910882	4.910931082	4.910931082
AA58794	3	2D Marker adult vs. fetal	0.857314271	4.809250141	4.448090919	3.267515415	0.565044105	0.573341181	0.49E-02
AA59178	3	2D Marker adult vs. fetal	1.9875003842	1.072838597	3.616532631	3.470451129	1.07047288	0.570591633	0.333930237
R81247	3	2D Marker adult vs. fetal	9.2855533478	4.4465905	3.6072685912	3.247665912	16.08923595	0.545116894	0.620081258
W8112	3	2D Marker adult vs. fetal	0.702500778	3.262302699	3.262302699	3.262302699	0.343082177	0.343082177	0.343082177
AA45485	3	2D Marker adult vs. fetal	1.297530164	3.13037216	3.13037216	3.13037216	11.3003685	0.223635384	0.223635384
N67048	2	2D Marker adult vs. fetal	0.162201012	0.397763394	0.297763394	0.30307128	0.27650405	0.233201513	10.14341663
AA77855	2	2D Marker adult vs. fetal	1.0340778279	1.0340778279	1.167591655	0.982841463	0.803409454	0.803409454	0.77581752
H51117	2	2D Marker adult vs. fetal	0.537673252	0.6873838102	0.5746220043	0.730285971	0.685326033	0.743516336	0.396538923
AA31174	5	2D Marker adult vs. fetal	0.62191585	0.62191585	0.62191585	0.62191585	0.4863862	0.4863862	0.4863862
AA77137	2	2D Marker adult vs. fetal	1.5231155704	1.0779766901	2.050526495	1.946322765	2.357520694	1.175923154	1.202805263
R05561	0	2D Marker adult vs. fetal	6.880519553	3.784035355	5.786261707	4.735376832	4.524485682	5.169772778	22.0681959
R16849	2	2D Marker adult vs. fetal	1.239798305	1.31798538	1.889965808	0.6639751034	1.415959831	1.415959831	7.595930359
AAB8167	2	2D Marker adult vs. fetal	1.2171209137	0.760767768	0.8461452007	1.070284972	1.1026569725	1.097445419	4.746402103
AA10115	2	2D Marker adult vs. fetal	1.5882989813	2.638659189	2.650975832	2.650975832	0.848297875	0.848297875	0.848297875
AA486822	2	2D Marker adult vs. fetal	1.638331704	2.664659174	2.616207444	2.282638599	1.570776152	1.570776152	1.570776152
AA69247	2	2D Marker adult vs. fetal	2.7105734034	2.7060573403	2.7060573403	2.7060573403	1.978934745	2.016294545	2.016294545
AA04049	3	2D Marker adult vs. fetal	6.383830468	5.671979456	5.681374244	4.838768605	9.255551514	9.255551514	9.255551514
AA22707	2	2D Marker adult vs. fetal	1.6822629404	2.164212705	2.1156465285	2.1051975731	2.9252853019	2.1214107168	2.891292998
NG2170	15	2D Marker adult vs. fetal	33.52697234	33.52697234	33.52697234	29.212938971	31.571024738	31.583171792	31.583171792
N27010	10	2D Marker adult vs. fetal	9.750528346	7.5309528346	7.5309528346	12.652020531	12.652020531	12.652020531	12.652020531
AA44185	16	2D Marker adult vs. fetal	5.25772564	5.25772564	5.25772564	7.014687482	7.014687482	7.014687482	7.014687482
AA45924	18	2D Marker adult vs. fetal	10.419706808	7.984616838	11.216572820	11.216572820	9.02612452	9.02612452	9.02612452
N7883	16	2D Marker adult vs. fetal	7.68464154	5.435265693	7.613477118	7.803876858	8.882600715	8.882600715	8.882600715
AA28719	18	2D Marker adult vs. fetal	8.071113381	20.07265983	19.16842124	13.654787922	27.057383931	31.571024738	31.583171792
AA47375	16	2D Marker adult vs. fetal	10.029716227	10.029716227	11.216572820	11.216572820	11.216572820	11.216572820	11.216572820
AA45351	7	2D Marker adult vs. fetal	3.30738388	3.622685625	3.622685625	3.622685625	3.670318593	4.776265186	4.776265186
AA05834	16	2D Marker adult vs. fetal	7.542505447	11.216572820	11.216572820	11.216572820	30.053197817	33.053197817	33.053197817
AA22789	17	2D Marker adult vs. fetal	13.32651569	9.041955128	16.97842166	12.44208045	15.162802039	28.234782305	30.146052309
AA45813	15	2D Marker adult vs. fetal	23.520146462	18.503051373	18.503051373	27.057383931	31.81897761	31.81897761	31.81897761
AA03755	16	2D Marker adult vs. fetal	10.029716227	10.029716227	12.727353005	12.727353005	12.727353005	12.727353005	12.727353005
AA45351	7	2D Marker adult vs. fetal	34.777842058	35.122567582	35.122567582	35.122567582	35.122567582	35.122567582	35.122567582
AA18133	15	2D Marker adult vs. fetal	27.207120577	18.081706489	27.212120505	29.689581169	29.689581169	30.053197817	30.053197817
AC22410	15	2D Marker adult vs. fetal	31.309593943	29.352102192	30.350704958	35.089758599	38.06106397	27.05519888	27.05519888
AA45243	15	2D Marker adult vs. fetal	27.208357468	24.45751672	26.021072	25.513173383	28.30704958	19.80388123	19.80388123
AA45245	15	2D Marker adult vs. fetal	12.54762444	10.029716227	12.54762444	12.54762444	13.803671361	17.939257323	17.939257323
AA32301	15	2D Marker adult vs. fetal	34.777842058	35.122567582	35.122567582	35.122567582	35.122567582	35.122567582	35.122567582
AA77334	15	2D Marker adult vs. fetal	10.419706808	7.984616838	11.216572820	11.216572820	11.216572820	11.216572820	11.216572820
AA22410	15	2D Marker adult vs. fetal	64.2525491623	43.912607527	58.30704958	58.30704958	58.30704958	58.30704958	58.30704958
AA45245	15	2D Marker adult vs. fetal	78.7226545	67.8571628	72.021072	72.021072	72.021072	72.021072	72.021072
AA45245	16	2D Marker adult vs. fetal	12.54762444	10.029716227	12.54762444	12.54762444	13.803671361	17.939257323	17.939257323
AA37651	23	2D Marker adult vs. fetal	13.60023495	12.18127765	12.18127765	12.18127765	12.18127765	12.18127765	12.18127765
AA45923	23	2D Marker adult vs. fetal	13.60023495	12.18127765	12.18127765	12.18127765	12.18127765	12.18127765	12.18127765
AA45923	0	2D Marker adult vs. fetal	10.419706808	7.984616838	11.216572820	11.216572820	11.216572820	11.216572820	11.216572820
N78821	17	2D Marker adult vs. fetal	18.16280113	18.16280113	18.16280113	18.16280113	18.16280113	18.16280113	18.16280113
AA20109	21	2D Marker adult vs. fetal	16.15200919	12.25195771	19.84085164	19.84085164	19.84085164	19.84085164	19.84085164
AA45245	17	2D Marker adult vs. fetal	18.16280113	18.16280113	18.16280113	18.16280113	18.16280113	18.16280113	18.16280113
AA15047	21	2D Marker adult vs. fetal	15.19512412	13.60023495	15.19512412	13.60023495	13.60023495	13.60023495	13.60023495
AA45923	23	2D Marker adult vs. fetal	17.24240499	13.60023495	17.24240499	13.60023495	13.60023495	13.60023495	13.60023495
AA45923	0	2D Marker adult vs. fetal	10.419706808	7.984616838	11.216572820	11.216572820	11.216572820	11.216572820	11.216572820
AA22427	21	2D Marker adult vs. fetal	17.24240499	13.60023495	17.24240499	13.60023495	13.60023495	13.60023495	13.60023495
AA77265	23	2D Marker adult vs. fetal	13.60023495	12.18127765	12.18127765	12.18127765	12.18127765	12.18127765	12.18127765
AA45923	23	2D Marker adult vs. fetal	13.60023495	12.18127765	12.18127765	12.18127765	12.18127765	12.18127765	12.18127765
AA77265	21	2D Marker adult vs. fetal	17.24240499	13.60023495	17.24240499	13.60023495	13.60023495	13.60023495	13.60023495
AA77265	17	2D Marker adult vs. fetal	13.60023495	12.18127765	13.60023495	12.18127765	12.18127765	12.18127765	12.18127765
AA45923	21	2D Marker adult vs. fetal	13.60023495	12.18127765	13.60023495	12.18127765	12.18127765	12.18127765	12.18127765
AA45923	0	2D Marker adult vs. fetal	10.419706808	7.984616838	11.216572820	11.216572820	11.216572820	11.216572820	11.216572820
AA45923	21	2D Marker adult vs. fetal	17.24240499	13.60023495	17.24240499	13.60023495	13.60023495	13.60023495	13.60023495
AA45923	23	2D Marker adult vs. fetal	13.60023495	12.18127765	13.60023495	12.18127765	12.18127765	12.18127765	12.18127765
AA45923	0	2D Marker adult vs. fetal	10.419706808	7.984616838	11.216572820	11.216572820	11.216572820	11.216572820	11.216572820
AA45923	21	2D Marker adult vs. fetal	17.24240499	13.60023495	17.24240499	13.60023495	13.60023495	13.60023495	13.60023495
AA45923	23	2D Marker adult vs. fetal	13.60023495	12.18127765	13.60023495	12.18127765	12.18127765	12.18127765	12.18127765
AA45923	0	2D Marker adult vs. fetal	10.419706808	7.984616838	11.216572820	11.216572820	11.216572820	11.216572820	11.216572820
AA45923	21	2D Marker adult vs. fetal	17.24240499	13.60023495	17.24240499	13.60023495	13.60023495	13.60023495	13.60023495
AA45923	23	2D Marker adult vs. fetal	13.60023495	12.18127765	13.60023495	12.18127765	12.18127765	12.18127765	12.18127765
AA45923	0	2D Marker adult vs. fetal	10.419706808	7.984616838	11.216572820	11.216572820	11.216572820	11.216572820	11.216572820
AA45923	21	2D Marker adult vs. fetal	17.24240499	13.60023495	17.24240499	13.60023495	13.60023495	13.60023495	13.60023495
AA45923	23	2D Marker adult vs. fetal	13.60023495	12.18127765	13.60023495	12.18127765	12.18127765	12.18127765	12.18127765
AA45923	0	2D Marker adult vs. fetal	10.419706808	7.984616838	11.216572820	11.216572820	11.216572820	11.216572820	11.216572820
AA45923	21	2D Marker adult vs. fetal	17.24240499	13.60023495	17.24240499	13.60023495	13.60023495	13.60023495	13.60023495
AA45923	23	2D Marker adult vs. fetal	13.60023495	12.18127765	13.6002				

W75892	19	2D Marker adult vs. total	53.6202013	39.30704958	56.67674826	53.8292161	46.2598911	20.27191765	17.02340594	12.4286249	13.0397038	19.7828359	
W70734	19	2D Marker adult vs. total	50.1395524	37.1918634	47.5751429	45.25023946	16.00676331	13.02269894	13.7703984	19.3128055	19.12043735	10.53049868	
H57136	13	2D Marker adult vs. total	10.27198246	9.124677788	11.15252179	10.05959216	23.82885337	23.14775465	17.64178704	22.6131072	27.82013988	19.25421118	
AA709414	23	2D Marker adult vs. total	72.8121734	37.16164602	36.30704958	68.6734991	62.23232659	54.4411949	29.75702023	29.71152223	30.84739122	35.55131266	37.82023417
W65401	7	2D Marker adult vs. total	5.747852929	5.328867725	6.071926118	6.40146448	5.626562584	5.582316414	1.032286392	1.038420135	1.71624513	1.31035945	2.22330134
AA45554	15	2D Marker adult vs. total	23.0291313	15.24754038	25.737051515	22.61565934	19.283691141	19.14163801	9.283691141	10.0106874	10.241574	10.0915821	5.46295703
AA428042	9	2D Marker adult vs. total	8.022234644	7.31169895	8.025622777	9.479217729	8.6254856534	7.544637481	50.76520202	25.54849687	20.5627043	18.97220626	14.02651814
AA427725	9	2D Marker adult vs. total	12.65284684	13.073265112	4.261153234	12.4290971	12.0148332	24.1811519	28.92515328	23.95557184	27.67022775	19.53259151	19.26976868
N51280	9	2D Marker adult vs. total	6.623233815	5.874959304	7.385168272	7.520693106	7.757574650	8.161620093	9.865251	31.8167829	22.571892	33.55516319	0.272525082
AA281347	8	2D Marker adult vs. total	5.089786513	4.840487592	4.841559344	4.84251025	4.490660004	4.492624512	22.81705154	14.8674075	16.0871733	26.21817205	21.27330752
AA412960	8	2D Marker adult vs. total	4.965180521	4.501823294	4.859223292	4.860365956	4.480263951	24.41674925	26.1691313	18.18713838	24.41674925	22.8465167	22.8465167
N98485	9	2D Marker adult vs. total	7.125465927	6.544129851	7.7771664	7.556752859	8.476056808	6.411668318	21.24752903	20.0377128	19.99045362	15.28211832	16.22577894
AA489209	7	2D Marker adult vs. total	5.051914729	5.0224950	4.8464705412	6.046521655	6.650662914	4.920069122	3.0356518537	2.9156518696	1.658183461	1.10759793	2.259717208
W61581	9	2D Marker adult vs. total	7.394753912	7.353526925	11.169355158	8.81000227	9.21761776	6.887179584	34.10197475	31.17716156	23.0122251	24.42864307	49.3715184
N51018	7	2D Marker adult vs. total	5.68510102	38.3926305	6.032538458	5.21842942	6.28226452	17.3222468	0.895593739	0.822653024	6.791305055	6.586840211	0.397310326
AA465201	7	2D Marker adult vs. total	6.950622778	6.952850011	7.93285008	7.93285008	7.93285008	7.93285008	7.93285008	7.93285008	7.93285008	7.93285008	6.59558359
W65471	7	2D Marker adult vs. total	7.215222797	6.544129851	8.025622381	8.167656349	8.167656349	12.55657547	21.76146549	22.52871009	1.889011302	4.29885207	3.911602
AA446521	7	2D Marker adult vs. total	6.616843406	3.668842167	10.1148224	5.389292381	5.404776773	6.067786717	1.03411616	1.422816774	15.19320754	7.416723464	1.40456398
AA465082	7	2D Marker adult vs. total	6.961269598	6.280975268	6.4414649774	7.072922475	7.565398009	6.402493823	5.030129346	5.030129346	1.456285641	2.37935622	4.17921622
AA44295	7	2D Marker adult vs. total	10.535688687	6.935251212	8.025620502	8.7150104314	8.7150104314	8.7150104314	8.7150104314	8.7150104314	8.658189404	2.325607801	3.97310326
N90003	7	2D Marker adult vs. total	6.996205007	7.777346527	5.122405597	7.694762829	7.694762829	8.111035017	8.111035017	8.111035017	3.214957653	5.781751821	6.1678428
AA009284	7	2D Marker adult vs. total	6.203527731	6.725228727	5.932223216	5.932223216	5.932223216	6.647324656	6.220425252	2.865921256	1.154565455	1.284565455	0.885776492
A1015368	12	2D Marker adult vs. total	4.73931812	5.632137227	6.299132956	4.5116454865	31.93521371467	32.51612495	19.81340558	22.2287943	29.07922706	24.0952182	25.86570587
AA478268	6	2D Marker adult vs. total	6.457623065	4.2186568521	6.072467356	5.821054619	5.705783899	4.27313179	11.9615041	10.98922076	11.69181426	14.11437595	16.20593802
AA608583	13	2D Marker adult vs. total	8.710219349	6.632651914	10.892670508	6.897784524	7.755924785	8.29174413	4.615478182	4.829174413	3.2565727167	3.658189404	4.180851252
AA505045	13	2D Marker adult vs. total	10.86563107	9.677783398	9.120702321	9.862800468	10.0505524	6.647324656	2.039052485	2.039052485	2.039052485	2.039052485	2.039052485
AA407693	13	2D Marker adult vs. total	6.175926959	7.7774740452	10.846325051	7.632223468	5.432223468	5.432223468	23.53565098	22.050453745	23.183942426	18.65535098	3.007772428
A202226	12	2D Marker adult vs. total	6.729813291	7.048662519	10.38317033	8.186857819	8.677781639	6.5456271414	20.27073033	18.6168657678	17.23229505	24.007151361	28.53431351
H67106	9	2D Marker adult vs. total	6.797311755	6.4959379167	6.699757753	6.245157348	6.155332271	7.944253554	48.602767655	42.396577553	23.161776775	29.1637751	30.15231052
WB450	12	2D Marker adult vs. total	8.323304604	8.323304604	8.323304604	8.323304604	8.323304604	8.323304604	8.323304604	8.323304604	8.323304604	24.56723452	24.56723452
NS331	13	2D Marker adult vs. total	12.01197758	0.755232839	1.042455794	1.042455794	1.042455794	1.042455794	1.042455794	1.042455794	1.042455794	1.042455794	1.042455794
AA405000	12	2D Marker adult vs. total	6.121595847	6.2295232058	7.7215050548	7.171810658	7.171810658	7.171810658	25.4356516191	26.80516179	23.914731213	20.13761267	22.51685956
T51539	12	2D Marker adult vs. total	6.489504146	6.570555251	6.489504146	6.489504146	6.489504146	6.489504146	6.489504146	6.489504146	6.489504146	37.1324121	54.37165159
NS9764	12	2D Marker adult vs. total	8.511173168	8.303765658	8.303765658	8.303765658	8.303765658	8.303765658	8.303765658	8.303765658	8.303765658	21.26531459	22.76220205
AA521546	11	2D Marker adult vs. total	18.0187483	11.98813991	20.04539147	19.7217852	19.7217852	19.7217852	23.076547267	23.076547267	25.614582767	25.614582767	21.762020578
AA42851	11	2D Marker adult vs. total	11.81370828	9.1323244514	12.530771771	12.530771771	12.530771771	12.530771771	12.530771771	12.530771771	12.530771771	28.030415188	23.039582168
AA489383	11	2D Marker adult vs. total	10.07168453	6.407010185	6.407010185	6.407010185	6.407010185	6.407010185	6.407010185	6.407010185	6.407010185	15.48920559	15.48920559
AA490172	11	2D Marker adult vs. total	8.374768147	1.415297225	1.415297225	1.415297225	1.415297225	1.415297225	1.415297225	1.415297225	1.415297225	37.1324121	24.37165159
AA504477	12	2D Marker adult vs. total	5.417691216	3.050554541	5.417691216	5.417691216	5.417691216	5.417691216	5.417691216	5.417691216	5.417691216	19.19137432	15.10292793